

BIOLOGICAL ACTIVITY OF ESSENTIAL OILS FROM LEAVES AND FRUITS OF PEPPER TREE (*Schinus molle* L.) TO CONTROL RICE WEEVIL (*Sitophilus oryzae* L.)

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ABSTRACT

Rice weevil (*Sitophilus oryzae* L.) is a primary insect pest of stored grain. The development of resistance resulted in the application of synthetic insecticides. In recent years many plant essential oils have provided potential alternatives to currently used insect control agents. The Brazilian pepper tree (*Schinus molle* L. var. *areira* (L.) DC.) (Anacardiaceae) has different biological properties such as insecticidal activity. In this study, repellent, fumigant activity, nutritional indices, and feeding deterrent action were evaluated on *S. oryzae* adults. Filter paper impregnation was used to test fumigant toxicity, whereas treated whole wheat was used to evaluate repellent activity and a flour disk bioassay was done to evaluate feeding deterrent action and nutritional index alteration. Leaf essential oils showed repellent effects at both concentrations (0.04 and 0.4% w/w), while fruit essential oils lacked repellent activity. Both plant oils altered nutritional indices. Fruit essential oils had a strong feeding deterrent action (62%) while leaves had a slight effect (40.6%). With respect to fumigant activity, neither of the essential oils was found to be toxic.

Key words: *Schinus molle*, *Sitophilus oryzae*, repellency, fumigant toxicity, nutritional indices, feeding deterrence.

INTRODUCTION

Harvest grains are basic human food products (Padín *et al.*, 2002). The presence of pests constitutes a serious on-going problem in stocking grains and its derived industry (Pérez Mendoza *et al.*, 2004).

Worldwide, between 5 and 15% of the total weight of cereals, oil plants, and legumes are lost after harvesting (Anonymous, 1989), and between 5 and 10% of these losses are due to the presence of pests (Hill, 1990). In Argentina, the losses caused by insects and/or mites are estimated between 7 and 10% of total production (Viale, 1995). The legislation of this country establishes the rejection of any merchandise with a single insect and/or live mite in any commercialization stage (Resolution N° 1975/94, Secretaría de Agricultura, Ganadería y Pesca).

The rice weevil (*Sitophilus oryzae* L.) is one of the pests of primary infestation of stored grains, widely spread worldwide, and very destructive. Because of its high incidence, synthetic insecticides have been used to control it.

Resistance and toxicity problems of the synthetic insecticides have resulted in the necessity of finding more effective and healthier alternatives. Thus, essential oils are the most tested products presently (Papachristos and Stamopoulos, 2002; Umoetok and Gerard, 2003; Zhang *et al.*, 2004; Tapondjou *et al.*, 2005; Ferrero *et al.*, 2006; Sánchez Chopa *et al.*, 2006, Stefanazzi *et al.*, 2006, Wang *et al.*, 2006). Different biological activities of plant derivatives have been demonstrated for the control of stored grain pests (Golob *et al.*, 1999; Rajendran and Sriranjini, 2008).

The Brazilian pepper tree (*Schinus molle* L. var. *areira* (L.) DC.) is associated with agricultural crops in boundaries, windbreaks, riverbank protection, and watershed conservation. In popular medicine, it is used for its astringent, diuretic, antispasmodic properties among others. In pest control, it is used as a fumigant, repellent, and ovicide (Ruffínengo *et al.*, 2005; Ferrero *et al.*, 2006).

The objective of the present study was to evaluate the fumigant, repellent and/or attractant, feeding deterrent activity, and the alterations in the nutritional physiology produced by the essential oils from leaves and fruits of *S. molle* var. *areira* on 3- to 4-day-old *S. oryzae* adults.

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MATERIALS AND METHODS

Insects

The insects came from a susceptible colony maintained in the Cátedra de Zoología Agrícola de la Universidad Nacional de Buenos Aires (UBA). They were bred in glass containers, 8 cm in diameter, 13 cm high, sealed with fine netting, maintained at 28 ± 1 °C, 60-70% HR, a 12:12 photoperiod, and using whole wheat grains as food.

Vegetal material

The vegetal organs (leaves and mature fruits) of the pepper tree were collected during the summer period in Bahía Blanca city, Buenos Aires, Argentina ($38^{\circ}41'$ S; $62^{\circ}17'$ W). A specimen was deposited in the Herbarium of the Departamento de Biología, Bioquímica y Farmacia de la Universidad Nacional del Sur (BBB) and identified as number CV 10444. The essential oils were isolated using fresh vegetal material by accelerated distillation of water vapor using a Clevenger-type device during 3 to 4 h, and were analyzed with gas chromatography and mass spectrometry (GC-MS HP5972A) in a HP5 (30 m x 0.25 mm) column with a temperature program of 50 °C during 2 min, a 5 °C min⁻¹ ramp, and a final temperature of 200 °C during 15 min. The composition of the oils from both vegetal organs can be observed in Table 1. Oil yield was 0.22% and 0.42% (w/v) for leaves and fruits, respectively. The yield was calculated using fresh vegetal material.

Bioassays

Repellent activity of the essential oils from leaves and fruits of *S. molle* var. *areira* on *S. oryzae* adults. An experimental field made of a central box connected by cylinders to four boxes symmetrically distributed around the first was used. Two boxes received 2 g of whole wheat treated with 2 mL of hexanic oil solutions at 0.04 and 0.4% (w/v), and 2 g rations impregnated with 2 mL of hexane were put in the remaining boxes as controls. In the central box, 40 unsexed 3 to 4 days old adults were released. After 24 h, the preference index (PI) was calculated using the Equation [1].

$$PI = \frac{(\text{percentage of insects in treated ration}) - (\text{percentage of insects in non-treated ration})}{(\text{percentage of insects in treated ration}) + (\text{percentage of insects in non-treated ration})} \quad [1]$$

where PI: -1.00 to -0.10 indicates repellent plant, PI: -0.10 to +0.10 neutral plant, and PI: +0.10 to +1.00 attractant plant. Furthermore, the repellency percentage was calculated based on the number of insects found in the control at the end of 24 h (Procopio *et al.*, 2003). Three replicates were carried out.

Table 1. Essential oil constituents from leaves and fruits of *Schinus molle* var. *areira*.

Compound	Leaves	Fruits
	%	
Limonene	15.68	40.34
α -Phellandrene	13.80	24.47
Elemol	9.00	
β -Cubebene	7.30	
Camphene	5.31	1.19
δ -Cadinene	5.26	
γ -Eudesmol	3.61	
α -Pinene	3.56	2.96
β -Eudesmol	2.80	
β -Pinene	2.13	0.58
β -Myrcene	1.72	16.35
Sabinene	1.48	
Caryophyllene	1.36	1.61
Triciclene	0.82	
Bornyl acetate	0.82	
1-Terpinen-4-ol		3.34
β -Phellandrene		2.39
3-Carene		1.35
Methyl octanoate		1.32
2-Carene		0.47
α -Humulene		0.25
Copaene		0.10

Nutritional indices and antifeeding activity of the essential oils from leaves and fruits of *S. molle* on *S. oryzae* adults. To evaluate the antifeeding activity and the alteration in the nutritional physiology in adults, 1.6 cm diameter disks of wheat flour were prepared (Huang *et al.*, 2002). Aliquots of 200 μ L from a flour suspension in water (10 g in 50 mL) were put on plastic dishes to form the disks and were left all night to dry in a chamber at 25 °C temperature and 60 to 70% relative humidity (Dalvo, model MCI/2 V.c.a 220, Argentina). The disks were weighed registering values between 70 to 78 mg. Hexanic solutions of the oils were prepared at concentrations of 0, 0.5, 1, 2, and 4 mg disk⁻¹. Two disks of wheat flour were impregnated with 5 μ L of these solutions, weighed, and put in separated containers. A control group with hexane treated disks was prepared. Ten 3- to 4-days-old adult insects previously weighed on a scale (FX/FY series FX400, Frankfurt, Germany) were put into each container. After maintaining them during 72 h in controlled conditions, the weight of the disks, mortality, and weight of insects alive were registered. Six replicates were carried out. Each assay was repeated independently at least three times. The nutritional indices were calculated:

Relative growth rate (RGR) = $(A-B)/(B \times \text{day})$, where A = weight of insects alive on the third day/number of insects alive on the third day; B = original weight of insects/total number of insects; relative consumption rate (RCR) which indicates the consumption of the insects related to their initial weight and the duration of the assay $RCR = D/(B \times \text{day})$, where D = biomass ingested (mg)/number of insects alive on the third day; efficiency of conversion of ingested food (ECI) (%) that indicates the quantity of food used for weight gain in the insects, $ECI = (RGR/RCR) \times 100$. To obtain the antifeeding effect (AE), $EA (\%) = [(C-T)/C] \times 100$ was calculated where C = consumption of control disks (mg) and T = consumption of treated disks (mg) (Farrar *et al.*, 1989).

Fumigant activity of the essential oils from leaves and fruits of *S. molle* var. *areira* on *S. oryzae* adults. Filter papers, 7 cm in diameter, were impregnated with 1 mL of hexanic solutions of the essential oils from leaves and fruits at concentrations of 10, 20, 40, and 80 mg L⁻¹. Hexane was used as a control. The solvent was left to evaporate during 5 min. The treated filter paper was put at the bottom of a 350 mL glass jar. Ten insects were put in each small glass tube of 5 cm high and 3 cm in diameter, with whole wheat and both open ends, covered with a fine netting to avoid them to scape. Each vial was suspended with a metal thread in the geometrical center of the jar which was hermetically sealed with a cover. The assay was carried out at controlled temperature and relative humidity conditions (28 ± 1 °C, 60-70% RH), and a 12:12 photoperiod. Three replicates were done. Each experiment was repeated independently at least three times. The percentage of mortality was evaluated after 72 h and expressed in mg L⁻¹ of air.

Statistical analysis of data

The data were analyzed by simple ANOVA test, completely randomized design prior normalization with $(\sqrt{x+1})^{1/2}$, using the replicates corresponding to each assay, and the means were separated using the minimum significant differences test with Microsoft Excel (MST, $p \leq 0.05$) (Zar, 1999).

RESULTS

Repellent activity of the essential oils from leaves and fruits of *S. molle* var. *areira* on *S. oryzae* adults. The repellent and/or attractant effects produced by the essential oils of *S. molle* on *S. oryzae* adults can be observed in Table 2.

It can be inferred from the analysis of Table 2 that the oil from leaves at both concentrations produced a repellent effect. However, the oil from fruits at the highest concentration produced an attractant effect whereas the lowest did not show any.

Table 2. Repellent activity of essential oil from leaves and fruits of the Brazilian pepper tree (*Schinus molle* var. *areira*) on adults rice weevils (*Sitophilus oryzae*).

Oil	Concentration	Repellency	PI
	% p/v	%	
Leaf	0.04	65.00	-0.30
	0.40	71.47	-0.40
Fruit	0.04	50.03	-0.01
	0.40	49.73	0.50

PI: Preference index: -1.0 to -0.1 repellent plant; -0.1 to +0.1 neutral plant, and +0.1 to +1.0 attractant plant.

Nutritional indices and antifeeding activity. In Tables 3 and 4, the results corresponding to the effects produced by both oils in relation to the nutritional indices and the antifeeding activity on *S. oryzae* adults are shown.

The oil from fruits modified the nutritional indices. RGR decreased in all the concentrations and highly significant differences were observed ($p \leq 0.01$) (F: 11.34, df: 20). Differences in RCR between the concentrations 0.5, 2, and 4 mg disk⁻¹ were observed with respect to the control ($p \leq 0.01$) (F: 6.2987, df: 20). ECI was only affected (at the highest) concentration ($p \leq 0.01$) (F: 6.6128, df: 20). A 54% mortality rate was observed at 4 mg disk⁻¹ concentration. AE (P = 0.1) (F: 2.2587, df: 16) of the oil from fruits of *S. molle* was slight at lower concentrations whereas it was high in the maximum concentration. Values greater than 51% were considered to be high.

The oil from leaves modified the nutritional indices, but contrary to what occurred with the oil from fruits, AE was slight ($p = 0.07$) (F: 2.733, df: 16). Furthermore, high significant differences were found ($p \leq 0.01$) (F: 4.3679, df: 20) in RGR at concentrations 1, 2, and 4 mg disk⁻¹ respect to the control. RCR decreased at maximum concentration respect to the control, and dropped off significantly at minimum concentration ($p \leq 0.05$) (F: 3.1604, df: 20). The same situation was observed with ECI ($p \leq 0.05$) (F: 3.3773, df: 20). A high mortality percentage was obtained with the essential oils from leaves at the highest concentration.

Fumigant activity of the essential oils from leaves and fruits of *S. molle* var. *areira* on 3- to 4-days-old *S. oryzae* adults. No fumigant toxicity was observed at the evaluated concentrations.

DISCUSSION

In this study, the essential oil of the leaves showed a repellent effect in both concentrations with repellency rates of 65 and 71.5% in the 0.04 and 0.4% concentrations,

Table 3. Nutritional indices and feeding deterrent activity of essential oil from fruits of Brazilian pepper tree (*Schinus molle* var. *areira*) on adults rice weevils (*Sitophilus oryzae*).

Concentration	RGR	RCR	ECI	Mortality	AE
mg disk ⁻¹	mg mg ⁻¹ d ⁻¹			%	
0	0.02c	0.28c	7.61b	2b	0
0.5	-0.07b	0.17ab	-47.12b	18b	30.68a
1	-0.06b	0.2bc	-71.96b	14b	20.11a
2	-0.08b	0.18b	-51.11b	14b	28.75a
4	-0.20a	0.09a	-312.2a	54a	62.07a

RGR: relative growth rate. RCR: relative consumption rate. ECI: efficiency conversion of ingested food. AE: antifeeding effect. Different letters in the same column indicate significant differences ($p \leq 0.05$) between treatments according to ANOVA and minimum significant difference (MSD).

Table 4. Nutritional indices and feeding deterrent activity of essential oil from leaves of Brazilian pepper tree (*Schinus molle* var. *areira*) on adults rice weevils (*Sitophilus oryzae*).

Concentration	RGR	RCR	ECI	Mortality	AE
mg disk ⁻¹	mg mg ⁻¹ d ⁻¹			%	
0	0.05c	0.3b	17.63c	2b	0
0.5	-0.009bc	0.28b	-8.86bc	12b	-9.84a
1	-0.08ab	0.22ab	-60.7abc	36a	18.36a
2	-0.1ab	0.2ab	-69.87ab	36a	24.63a
4	-0.14a	0.16a	-111.34a	58a	40.60a

RGR: relative growth rate. RCR: relative consumption rate. ECI: efficiency of conversion of ingested food. AE: antifeeding effect. Different letters in the same column indicate significant differences ($p \leq 0.05$) between treatments according to ANOVA and minimum significant difference (MSD).

respectively. Using essential oil from the leaves of *Artemisia princeps* Pamp. (Asteraceae), Liu *et al.* (2006) observed a good repellent activity on *S. oryzae*. On the other hand, Amelot *et al.* (2003) demonstrated that oils of *Ageratum conyzoides* L. (Asteraceae) leaves did not produce a repellent effect on this insect. The oil from fruits of *S. molle* var. *areira* resulted attractant only at the highest concentration. The observed differences in the effects produced by the essential oils could be due to the presence of different secondary metabolites in both vegetal organs (Murray *et al.*, 2005).

The AE of the oil from leaves of *S. molle* was slight, whereas with fruits it was high. Valera *et al.* (2003) found that the essential oil of *Coleus amboinicus* Loureiro (Lamiaceae) was not able to induce any response in the feeding behavior of *S. oryzae* at high doses (1.25%), there was not mortality during the 60 h of bioassay exposition.

Oils from leaves and fruits of *S. molle* modified the nutritional physiology of *S. oryzae*, altering RGR, RCR, and ECI. The oil from fruits, at maximum dose (4 mg disk⁻¹) produced high mortality which could be due to an inhibition in the feeding behavior (behavioral effect) based on the decrease of the RCR and the high antifeeding effect. The oil from leaves at maximum concentration also produced high mortality as a consequence of a possible

post-ingestion toxicity due to the slight AE and the decrease in ECI. Liu and Ho (1999) observed modifications in RGR, but neither in the consumption rate, nor in ECI with *Evodia rutaecarpa* (Rutaceae) in another *Sitophilus* species. The essential oil of *Tagetes ternifolia* (Asteraceae) also produced post-ingestion toxicity in other pest of stored grains *Tribolium castaneum* (Stefanazzi *et al.*, 2006).

The essential oils from leaves and fruits of *S. molle* did not produce fumigant activity on *S. oryzae* adults. The lack of this activity could be due to the fact that the entrance of these compounds is through the cuticle, to the need of higher concentrations to reach the mortality, or more exposition time. A similar situation was observed by Pascual Villalobos *et al.* (2004) using essential oils from leaves of *Ocimum basilicum* L. (Lamiaceae). Authors such as Negahban *et al.* (2007), Negahban and Moharrampour (2007) demonstrated fumigant activity with essential oils of *Artemisia* (Asteraceae) and *Eucalyptus* (Myrtaceae) species on *S. oryzae*. In *Sitophilus granarius*, oils of *Salvia hydrangea* DC., former Benth (Lamiaceae) produced toxicity (Kotan *et al.*, 2008).

CONCLUSIONS

This study demonstrated that the essential oil from leaves of *Schinus molle* var. *areira* produced repellency

on *S. oryzae*, while the fruit resulted attractant at 0.4%, and neutral at a concentration of 0.04%. Both oils altered nutritional physiology, showed antifeeding activity, and did not exhibit fumigant activity.

For all these reasons, we can infer that the essential oils of leaves and fruits of *Schinus molle* var. *areira* could be considered as a natural alternative in the control of *S. oryzae*.

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RESUMEN

Bioactividad de aceites esenciales de hojas y frutos del aguaribay (*Schinus molle* L.) en el gorgojo del arroz (*Sitophilus oryzae* L.). El gorgojo del arroz (*Sitophilus oryzae* L.) es un insecto-plaga de infestación primaria de granos. El uso de insecticidas sintéticos ha desarrollado fenómenos de resistencia. En los últimos años los aceites esenciales se presentan como una alternativa en el control de insectos-plaga. El aguaribay (*Schinus molle* L. var. *areira* (L.) DC.) (Anacardiaceae) es una planta con diferentes propiedades biológicas entre las que se destacan el uso como insecticida. El objetivo de este estudio fue evaluar la actividad fumigante, repelente, los índices nutricionales y la actividad antialimentaria de los aceites esenciales de hojas y frutos de *S. molle* var. *areira* en adultos de *S. oryzae*. Para la actividad fumigante se utilizó la técnica de impregnación de papeles de filtro; para la actividad repelente impregnación de trigo entero; y para los índices nutricionales y la actividad antialimentaria impregnación de discos de harina de trigo. El aceite esencial de hojas mostró efectos repelentes a ambas concentraciones (0,04 y 0,4% p/v), mientras que el de frutos no produjo repelencia. Ambos aceites alteraron la fisiología nutricional de *S. oryzae*. El aceite de frutos produjo un efecto antialimentario fuerte (62%) y el de hojas leve (40,6%). No se observó actividad fumigante.

Palabras clave: *Schinus molle*, *Sitophilus oryzae*, repelencia, toxicidad fumigante, índices nutricionales, efecto antialimentario.

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LIFE TABLE PARAMETERS AND CONSUMPTION RATE OF *Cydnodromus picanus* Ragusa, *Amblyseius graminis* Chant, AND *Galendromus occidentalis* (Nesbitt) ON AVOCADO RED MITE *Oligonychus yothersi* (McGregor) (ACARI: PHYTOSEIIDAE, TETRANYCHIDAE)

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ABSTRACT

The avocado red mite *Oligonychus yothersi* (McGregor) is the major leaf pest in Chile's avocado orchards. It affects leaf physiology and makes it necessary to seek new natural enemies to interact with low population densities of *O. yothersi*. The potentiality of three predator mites: *Cydnodromus picanus* Ragusa, *Amblyseius graminis* Chant, and *Galendromus occidentalis* (Nesbitt) was evaluated under laboratory conditions (27 ± 1.93 °C, $87 \pm 3.61\%$ H.R. and 16:8 (L:D) photoperiod) on avocado leaf disks *Persea americana* Mill. var. Hass ($\varnothing = 5$ cm) by separately feeding eggs, immature, and adult females of *O. yothersi*, and registering postembryonic development, consumption, as well as life table parameters. The postembryonic development of *C. picanus* was significantly lower (5.46 days) compared to both *A. graminis* (7.33 days) and *G. occidentalis* (8.69 days) which were fed with immature *O. yothersi*. The life table parameters of *C. picanus* were net reproductive rate $R_0 = 25.41$, finite rate of increase $\lambda = 1.29$, and mean generation time $T = 12.46$. The net intrinsic rate of increase (r_m) was significantly higher for *C. picanus* ($r_m = 0.25$) in contrast with *G. occidentalis* ($r_m = 0.19$), while *A. graminis* showed $r_m = -0.06$ indicating that its population didn't have descendants. Under laboratory conditions, r_m registered by *C. picanus* is an indicator of its predatory potential to control *O. yothersi*. It can be assumed that the pest population reduction pattern could be maintained under field conditions.

Key words: postembryonic development, predation, pollen, biological control.

INTRODUCTION

The avocado, *Persea americana* Mill. (Lauraceae), is the second most cultivated fruit tree in Chile after vineyards, and covers an area of 39 302.59 ha of which 56% is concentrated in the Valparaíso Region (INE, 2007). Furthermore, Chile is the second world exporter of avocados, mainly the Hass variety, with approximately 165 000 t exported during the 2006-2007 season (Comité de Paltas, 2007).

Nevertheless, there is an economic loss associated with exports because of the presence of pests such as *Pseudococcus longispinus* (Targioni & Tozzetti) (Hemiptera: Pseudococcidae), *P. calceolariae* (Maskell) (Hemiptera: Pseudococcidae), *Hemiberlesia lataniae*

(Signoret) (Hemiptera: Diaspididae), and *Heliothrips haemorrhoidalis* (Bouché) (Thysanoptera: Thripidae) (SAG, 2007). The most important economic avocado pest at a foliar level is *Oligonychus yothersi* (McGregor) (Acari: Tetranychidae) (Altieri and Rojas, 1999), commonly known as the avocado red mite, and var. Hass is the most susceptible to be attacked by this tetranychid. *Oligonychus yothersi* provokes a decrease in photosynthetic rate, stomatal conductance, and transpiration, negatively affecting the physiology of the avocado leaves (Schaffer *et al.*, 1986). This has a direct consequence on the quality of the fruit and crop yield (Palevsky *et al.*, 2007a), the same as for *O. perseae* Turtle, Baker and Abbatiello (Acari: Tetranychidae) found in California, USA (Kerguelen and Hoddle, 2000; Takano-Lee and Hoddle, 2002).

The natural enemies associated with *O. yothersi* in avocado orchards in the Province of Quillota are *Stethorus histrio* Chazeau (Coleoptera: Coccinellidae) and *Oligota pygmaea* Solier (Coleoptera: Staphylinidae), density-dependent generalist predators. Both coleoptera present natural colonization in the orchard only when the pest population increases (Obrycki and Kring, 1998; Kishimoto,

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2003) without exerting the necessary regulation to avoid damage produced by the red mite at the leaf physiological level. This makes it necessary to incorporate new predators to the system to interact with low *O. yothersi* population densities in the Chilean avocado orchards managed with biological control agents.

The most important predators of phytophagous mites in the world belong to the Phytoseiidae (Shrewsbury and Hardin, 2003) family which are easily adaptable to perturbed habitats and intensely managed as is the case of fruit orchards (Croft and Luh, 2004). The generalist species do not require large mite pest population densities to be established in an orchard, and migrate to other places through aerial dispersion if they lack prey (Colfer *et al.*, 2003; Tixier *et al.*, 2006). In the absence of phytophagous mites, the generalists have the capacity to use food alternatives such as pollen grains, fungi spores, insects in the immature stages, plant nectar, and exudates (Croft *et al.*, 2004; Nomikou *et al.*, 2005; Bouras and Papadoulis, 2005).

To include new natural enemies in a biological control system, it is fundamental to know their biological and ecological characteristics. The potential of the predators on their prey (De Vis *et al.*, 2006b) can be estimated through population models and the construction of life tables, thus obtaining data about survival, longevity, reproduction, and descendants of the arthropod populations (Yu *et al.*, 2005; Gabre *et al.*, 2005; Yang and Chi, 2006; Ozman-Sullivan, 2006; Ferrero *et al.*, 2007). Food quality has a great influence on the formulation of biological parameters since it is indispensable to recognize the predator's consumption in each stage of the pest in order to predict its effectiveness as a natural enemy (Kishimoto, 2003; Gotoh *et al.*, 2006; Collier *et al.*, 2007), and potential impact on the prey (Hosseini *et al.*, 2005).

This study evaluated the demographic parameters and the consumption of *Cydnodromus picanus*, *Amblyseius graminis*, and *Galendromus occidentalis* on distinct stages of *O. yothersi*, first under laboratory conditions to identify the red mite's potential predators which would eventually be included in integrated pest management plans.

MATERIALS AND METHODS

Species studied. Three phytoseiid species were selected and evaluated as potential predators of the avocado red mite based on biological and ecological characteristics. *Cydnodromus picanus* Ragusa (Parasitiforms: Phytoseiidae) is a type III generalist phytoseiid from the Pica zone (20°15' S; 69°20' W), Tarapacá Region (Ragusa *et al.*, 2000) which is able to

withstand great thermal oscillations during throughout the day with scarce environmental humidity, and survive food scarcity. *Amblyseius graminis* Chant (Parasitiforms: Phytoseiidae) is a type III generalist phytoseiid (Croft *et al.*, 2004) collected on redstem stork's bill (*Erodium cicutarium* (L.) L'Hér.) (Geraniaceae) in avocado orchards in the La Cruz zone (32°49' S; 71°17' W), Valparaíso Region. *Galendromus (Metaseiulus) occidentalis* (Nesbitt) (Parasitiforms: Phytoseiidae) is a type II specialist phytoseiid (Blackwood *et al.*, 2004) collected on walnut trees (*Juglans regia* L.) (Juglandaceae) (Ragusa and Vargas, 2002) in Los Andes locality (32°49' S; 70°35' W), Valparaíso Region, and is a known mite predator of the *Oligonychus* genus (Shrewsbury and Hardin, 2003).

Site and study materials. The life table and consumption assays were carried out in the laboratories of Instituto de Investigaciones Agropecuarias (INIA) La Cruz, Valparaíso Region, between January and September 2007. Using a data logger, the Petri dish micro-climatic conditions were registered inside the laboratory, thereby obtaining a temperature of 27 ± 1.93 °C, relative humidity of $87 \pm 3.61\%$, and a 16:8 (L:D) photoperiod for all the assays. These micro-climatic conditions were used to register the maximum biological potential of the predatory species since these are susceptible to low humidity in the egg stage (De Vis *et al.*, 2006a). The experimental observations were carried out every 24 h with a 40X stereoscopic magnifying glass (Zeiss, Stemi, Göttingen, Germany). An adhesive (Point sticken blue, Point Chile S.A.) was used to avoid the mites from escaping.

Breeding of the avocado red mite. *Oligonychus yothersi* were bred massively on avocado leaves var. Hass, with a modified methodology (Oliveira *et al.*, 2001) using plastic containers (29 x 7 x 39.5 cm) at a temperature of 27 ± 2 °C, relative humidity of $50 \pm 10\%$, and a 16:8 (L:D) photoperiod. The micro-climatic conditions were registered with a digital thermo-hygrometer.

Phytoseiid breeding. The three predatory species selected were obtained in the phytoseiid breeding room located in the INIA La Cruz facilities. Subsequently, gravid females of this species were moved to the assay laboratory where they were bred on avocado leaf disks var. Hass infested with *O. yothersi* inside plastic containers (57 x 42 x 19 cm) opened at the top and covered with muslin to avoid contamination of the predatory mite populations. The assays were carried out with eggs laid by the first-generation females.

Postembryonic development. Egg-adult development was determined for each species of phytoseiid. Thirty gravid females were taken from each species and each female was placed inside an avocado leaf disk var.

Hass ($\varnothing = 5$ cm) confined with adhesive (sticken). They were eliminated after 5 h, leaving 1 egg per disk (1 egg = 1 replicate), and registering the duration of each developmental stage of the phytoseiid through the exuvium. Longevity of unmated individuals was obtained by making available, on a daily basis, ten 24-h-old eggs, 10 mobile immature individuals (protonymphs and deutonymphs), and five *O. yothersi* adult females. Daily consumption was registered for each phytoseiid.

Avocado (*Persea americana* Mill.) (Lauraceae) var. Hass and hairy brassica (*Hirschfeldia incana* (L.) Lagr.-Foss.) (Brassicaceae) pollen was evaluated as alternative food to verify the survival of the species when facing a scarcity of prey. Daily, avocado var. Hass and *H. incana* pollen was provided separately by means of a fine brush, along with registering postembryonic development and predator longevity. Water was provided by cotton threads through a hole in the leaf for assays with pollen, as well as for those without food supply.

Fertility and longevity. Thirty females of known age were placed in avocado var. Hass leaf disks (1 female = 1 replicate), integrating a male for 24 h every 7 days. Each female was given 15 mobile immature *O. yothersi* (protonymphs and deutonymphs). The phytoseiid eggs were counted and eliminated, recording longevity, fecundity, and consumption of the gravid females. To obtain descendants and the proportion of sexes, 10 females were randomly selected from the previous 30. Thus, the eggs of each female were counted and deposited on 10 infested Petri dishes with all the *O. yothersi* stage, respectively, thus recording data about fertility and proportion of sexes for the females of each species.

Statistical analysis. A completely random design was applied with 30 replicates per experiment. Postembryonic development, longevity, and consumption data were transformed by $\sqrt{x + 0.5}$ (Steel and Torrie, 1985). Subsequently, ANOVA and Tukey test ($p < 0.05$) were applied to evaluate the influence of food on postembryonic development and phytoseiid consumption.

The following were the calculated life table parameters (SAS Institute, 2007): (1) Net reproductive rate, $R_0 = l_m m_x$, being the number of females that produce a female during a generation or during their lifespan (Rabinovich, 1980); (2) intrinsic rate of increase, r_m being the maximum exponential multiplication rate of a whole population, and calculated as $1 = l_m \exp(-r_m)$ (Birch, 1948); (3) finite rate of increase, $\lambda = \exp(r_m)$ being the number of females that produce one female per day (Birch, 1948); and (4) generation time, $T = \sum l_m m_x / R_0$ being the time that passes between first and next generation oviposition (Rabinovich, 1980).

The Jackknife nonparametric resampling method was used to compare the parameters of the life table between

species, estimating the mean, variance, and standard error (Meyer *et al.*, 1986; La Rossa and Kahn, 2003) with the LIFETABLES software, SAS (Maia *et al.*, 2000), and SAS® (SAS Institute, 2007). The biological parameters were subsequently compared with the Tukey test ($P < 0.05$).

RESULTS

The time of postembryonic development of *C. picanus* observed was less compared to the other two predatory species ($F = 134.54$, $df = 2$, $p < 0.01$) when fed mobile immature *O. yothersi* (Table 1). With regard to the longevity of phytoseiids fed with mobile immature *O. yothersi*, *C. picanus* showed a greater duration of the adult stage than *A. graminis* and *G. occidentalis*, thus indicating that the supply of *O. yothersi* protonymphs and deutonymphs had a positive influence on the postembryonic development of *C. picanus* ($F = 167.30$, $df = 2$, $p < 0.01$). In relation to the percentage of immature phytoseiids that developed to the adult stage, a survival rate of 100% was registered for *C. picanus*, 86% for *G. occidentalis*, and only 10% for *A. graminis*.

By feeding *O. yothersi* eggs, the postembryonic development of *C. picanus* and *A. graminis* increased with respect to the predators fed with immature red mites, whereas *G. occidentalis* only reached the larval stage. Furthermore, *C. picanus* showed a 13% survival rate and *A. graminis* 6.6% in the immature stage (Table 1). It was confirmed that in the immature stage, *C. picanus*, *A. graminis*, and *G. occidentalis* do not consume adult females of the avocado red mite (Table 1).

Using avocado var. Hass pollen, the duration of the postembryonic development was found to be shorter for *A. graminis* than *C. picanus* ($F = 27.55$, $df = 1$, $p < 0.01$). Regarding longevity of the evaluated species, *A. graminis* individuals were significantly more long-lived ($F = 148.18$, $df = 1$, $p < 0.0001$) than *C. picanus*. Survival of immature phytoseiids that reached the adult stage was not significantly different between *A. graminis* (66.6%) and *C. picanus* (43.3%) ($F = 3.38$, $df = 1$, $p = 0.0713$), though *G. occidentalis* did not consume pollen and only developed to the larval stage (Table 2).

Using *H. incana* pollen, egg-adult development was observed to be less for *A. graminis* than *C. picanus* ($F = 177.21$, $df = 1$, $p < 0.01$), although longevity was significantly greater for *A. graminis* ($F = 345.48$, $df = 1$, $p < 0.0001$) than *C. picanus*. Furthermore, *A. graminis* showed a 60% survival rate of individuals in the immature stage that developed into the adult stage, whereas *C. picanus* registered a statistically similar 46.6% ($F = 1.05$, $df = 1$, $p = 0.3087$) (Table 2).

In terms of *C. picanus* longevity, a significant difference was obtained for the individuals fed with mobile immature

Table 1. Duration of postembryonic development and longevity (in days) of *Cydnodromus picanus*, *Amblyseius graminis*, and *Galendromus occidentalis* fed with *Oligonychus yothersi* in different stages.

Daily diet	State	Phytoseiid species (mean ± SE)					
		n	<i>C. picanus</i>	n	<i>A. graminis</i>	n	<i>G. occidentalis</i>
10 mobile immature <i>O. yothersi</i>	Egg	30	2.00 ± 0.00b	30	2.00 ± 0.00b	30	3.00 ± 0.00a
	Larva	30	1.00 ± 0.00b	30	1.00 ± 0.00b	30	2.10 ± 0.00a
	Protonymph	30	1.03 ± 0.03b	28	2.07 ± 0.18a	26	1.69 ± 0.09a
	Deutonymph	30	1.60 ± 0.09b	15	2.60 ± 0.13a	26	1.92 ± 0.10b
	Egg-Adult	30	5.46 ± 0.10c	3	7.33 ± 0.33b	26	8.69 ± 0.16a
	Adult (longevity)	30	60.03 ± 1.54a	3	18.00 ± 2.08c	26	29.08 ± 1.17b
10 <i>O. yothersi</i> eggs	Egg	30	2.00 ± 0.00b	30	2.00 ± 0.00b	30	3.00 ± 0.00a
	Larva	30	1.00 ± 0.00b	30	1.00 ± 0.00b	30	2.07 ± 0.11a
	Protonymph	29	2.89 ± 0.12b	21	3.80 ± 0.29a	0	-
	Deutonymph	10	3.90 ± 0.31a	7	3.00 ± 0.30a	0	-
	Egg-Adult	4	9.09 ± 0.62a	2	9.50 ± 0.50a	0	-
	Adult (longevity)	4	19.75 ± 2.42a	2	9.50 ± 1.50b	0	-
5 <i>O. yothersi</i> adult females	Egg	30	2.00 ± 0.00b	30	2.00 ± 0.00b	30	3.00 ± 0.00a
	Larva	30	1.00 ± 0.00b	30	1.00 ± 0.00b	30	2.40 ± 0.09a
	Protonymph	30	4.66 ± 0.08a	8	2.68 ± 0.13b	0	-
	Deutonymph	0	-	0	-	0	-
	Egg-Adult	0	-	0	-	0	-
	Adult (longevity)	0	-	0	-	0	-

Values with distinct letters in the row indicate significant differences according to Tukey ($p < 0.05$).

SE: standard error. n: number of individuals.

red mites (60.03 days) as compared with administering an exclusive diet of avocado var. Hass pollen (40.46 días) ($F = 62.74$, $df = 1$, $p < 0.0001$) and *H. incana* (22.5 días) ($F = 251.41$, $df = 1$, $p < 0.0001$), thus indicating that these two latter diets are a feeding alternative when prey is scarce. On the other hand, *A. graminis* registered a significantly greater longevity when fed avocado var. Hass pollen (78.10 días) ($F = 91.36$, $df = 1$, $p < 0.10001$) and *H. incana* (84.94 días) ($F = 86.85$, $df = 1$, $p < 0.0001$) compared with feeding on mobile immature red mites (18 days) (Table 1, Table 2).

On a water diet, *C. picanus* and *A. graminis* developed up to the protonymph stage. In contrast, *G. occidentalis* only reached the larval stage. Furthermore, *C. picanus* showed a longer duration in the protonymph stage compared with *A. graminis* ($F = 1158.03$, $df = 1$, $p < 0.01$) (Table 2).

As for depredation on immature *O. yothersi*, *G. occidentalis* registered consumption of the avocado red mite in the larval stage although *C. picanus* and *A. graminis* did not present depredation in this stage ($F = 457.40$, $df = 2$, $p < 0.01$), indicating that *G. occidentalis* needs to be fed to continue its postembryonic development. On the other hand, *C. picanus* and *G. occidentalis* registered less depredation in the protonymph stage than *A. graminis* ($F = 32.58$, $df =$

2, $p < 0.01$), a behavior also observed in deutonymphs ($F = 13.77$, $df = 2$, $p < 0.01$). Nevertheless, unmated *C. picanus* adults showed a greater depredation rate compared with unmated *A. graminis* and *G. occidentalis* adults ($F = 71.96$, $df = 2$, $p < 0.01$) (Table 3).

A greater depredation rate of mated *A. graminis* females on immature *O. yothersi* was observed as compared with *C. picanus* and *G. occidentalis* ($F = 306.67$, $df = 2$, $p < 0.01$) (Table 3).

Life table parameters

Cydnodromus picanus females showed gradual mortality over time in contrast with *A. graminis* and *G. occidentalis* which concentrated almost 80% mortality in 7 days (Figure 1). Furthermore, greater longevity was noted for *A. graminis* (25.7 days) and *C. picanus* (25.43 days) females in contrast with *G. occidentalis* (22.56 days) ($F = 5.44$, $df = 2$, $p = 0.006$). The three survival curves recorded for the distinct species were type I, thus indicating that mortality was mainly concentrated in long-lived individuals (Rabinovich, 1980).

There is no significant difference in the oviposition rate between the evaluated phytoseiid species ($F = 1.47$, $df = 2$, $p = 0.236$) (Figure 2). Comparing female fertility, *C. picanus* had a higher value than *G. occidentalis* and

Table 2. Influence of diet on duration of postembryonic development and longevity (in days) of *Cydnodromus picanus*, *Amblyseius graminis*, and *Galendromus occidentalis*.

Daily diet	State	Phytoseiid species (mean ± SE)					
		S ¹ (%)	<i>C. picanus</i>	S (%)	<i>A. graminis</i>	S (%)	<i>G. occidentalis</i>
<i>Persea americana</i> var. Hass pollen	Egg	100a	2.00 ± 0.00b	100a	2.00 ± 0.00b	100a	3.00 ± 0.00a
	Larva	100a	1.00 ± 0.00b	100a	1.00 ± 0.00b	100a	1.13 ± 0.06a
	Protonymph	96.6a	4.37 ± 0.18a	66.6b	2.40 ± 0.13b	0c	-
	Deutonymph	73.3a	3.50 ± 0.20a	66.6a	2.40 ± 0.21b	0b	-
	Egg-Adult	43.3a	10.86 ± 0.52a	66.6a	7.80 ± 0.26b	0b	-
	Adult (longevity)	43.3a	40.46 ± 1.11b	66.6a	78.10 ± 2.37a	0b	-
<i>Hirschfeldia</i> <i>incana</i> pollen	Egg	100a	2.00 ± 0.00b	100a	2.00 ± 0.00b	100a	3.00 ± 0.00a
	Larva	100a	1.00 ± 0.00b	100a	1.00 ± 0.00b	100a	1.03 ± 0.03a
	Protonymph	93.3a	2.67 ± 0.14a	60b	1.00 ± 0.00b	0c	-
	Deutonymph	83.3a	3.04 ± 0.12a	60b	1.05 ± 0.05b	0c	-
	Egg-Adult	46.6a	8.68 ± 0.26a	60a	5.05 ± 0.05b	0b	-
	Adult (longevity)	46.6a	22.50 ± 0.97b	60a	84.94 ± 2.85a	0b	-
Water	Egg	100a	2 ± 0.0 b	100a	2.00 ± 0.00b	100a	3.00 ± 0.00a
	Larva	100a	1 ± 0.0 b	100a	1.00 ± 0.00b	100a	1.15 ± 0.06a
	Protonymph	100a	4.73 ± 0.08 a	66.6b	1.10 ± 0.06b	0c	-
	Deutonymph	-	-	-	-	-	-
	Egg-Adult	-	-	-	-	-	-
	Adult (longevity)	-	-	-	-	-	-

Distinct letters indicate significant differences according to Tukey ($p < 0.05$).

S¹: survival of individuals expressed in %. SE: standard error. n: number of individuals.

A. graminis whose eggs were almost entirely infertile (Figure 3).

The life table parameters of the three phytoseiids fed with immature *O. yotheresi* showed that *C. picanus* showed higher R_0 , r_m , and λ than *G. occidentalis* ($F = 233.58$, $df = 3$, $p < 0.0001$; $F = 2390.05$, $df = 3$, $p < 0.0001$; $F = 215.61$, $df = 3$, $p < 0.0001$; $F = 2127.12$, $df = 3$, $p < 0.0001$), whereas *A. graminis* revealed $R_0 = 0.27$ indicating that the population of this specie decreases over time (Table 4). The biological parameters of *C. picanus* show that the population grew 25.41 times in 12.46 days (T), and for each female of the actual generation there will be 25.41 females in the next generation. Furthermore, for each female present on a given day, there will be almost 1.29 (λ) females the next day. Therefore, at any particular point in time, the number of females in the *C. picanus* population will increase at such a rate that a population growth of 25% (r_m) is expected from one day to the next. Moreover, comparing R_0 , r_m , T, and λ of *C. picanus* with *O. yotheresi*, it is observed that only the latter attains a higher R_0 ($F = 233.58$, $df = 3$, $p < 0.0001$), while the predator registered higher r_m and λ . In addition, generation time was significantly lower for *C. picanus* ($F = 215.61$, $df = 3$, $p < 0.0001$) demonstrating that it multiplied more rapidly than the red mite population (Table 4).

DISCUSSION

Consumption records during the postembryonic development of the three phytoseiids in the *O. yotheresi* egg, immature, and adult female stages indicated that protonymphs and deutonymphs of the avocado red mite are differentially predated by *C. picanus*, *A. graminis*, and *G. occidentalis*. This influenced the depredation rate by, morphology, prey stage, and the predators' mouth parts (Croft *et al.*, 2004), since the integuments of *O. yotheresi* adult females are more difficult to penetrate than those of immature prey (Kishimoto y Takagi, 2001; Furuichi *et al.*, 2005).

Regarding consumption, Ragusa *et al.* (2000) established that *C. picanus* fed with *Tetranychus urticae* C.L. Koch (Tetranychidae) eggs reach the adult stage in approximately 4 days, demonstrating a positive influence of this prey in the development of the phytoseiid compared with *O. yotheresi*. This would be explained by *T. urticae* egg morphology: spherical and easy to handle by the phytoseiids. In contrast, *O. yotheresi* eggs adhere to the surface of the avocado leaf making it difficult to capture, and consequently less attractive as food (Vantornhout, 2006).

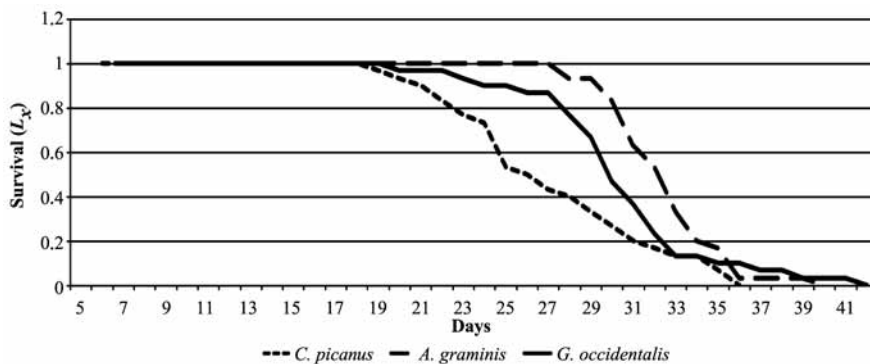
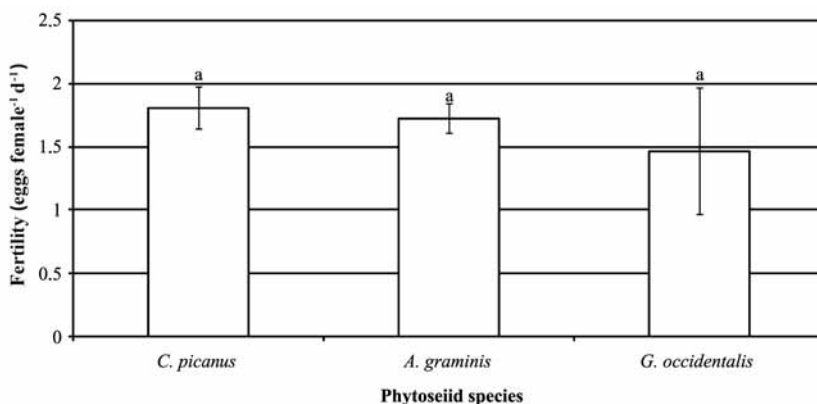


Figure 1. Survival curve of mated female *Cydnodromus picanus*, *Amblyseius graminis*, and *Galendromus occidentalis* fed with mobile immature *Oligonychus yothersi*.



Distinct letters indicate significant differences according to Tukey ($p < 0.05$).

Figure 2. Female fertility of *Cydnodromus picanus*, *Amblyseius graminis*, and *Galendromus occidentalis* fed with mobile immature *Oligonychus yothersi*.

The established classification with regard to alternative food was confirmed by pointing out *C. picanus* and *A. graminis* as type III generalists and *G. occidentalis* as a recognized type II specialist preferring the *Oligonychus* genus (Shrewsbury and Hardin, 2003; Croft *et al.*, 2004). Both generalists would be more adapted to conditions of food scarcity than *G. occidentalis* which need to feed on mites in order to develop. Ragusa *et al.* (2000) gave *Oxalis* sp. and *Ricinus* sp. pollen to *C. picanus* exhibiting survival rates of 52% and 44%, respectively in the immature stage. When fed with avocado var. Hass pollen, 43.3% of the population survived, converting it into an ideal alternative food in the absence of the red mite, and demonstrating another comparative advantage over *G. occidentalis*. It is also worth mentioning that *G. occidentalis* in commercial orchards is easily displaced by type III generalist phytoseiids. Slow and smaller-sized, it can also be transformed into prey for phytoseiids, and easily depredated by coleopters belonging to the *Stethorus*

and *Oligota* genera (Colfer *et al.*, 2003). Therefore, possible field releases of *G. occidentalis* could only be carried out when the *O. yothersi* population is high in the orchard, without being able to avoid the physiological damage provoked by the red mite on avocado leaves.

When *O. yothersi* population density is low and within a context of habitat management, it would be possible to carry out preventive releases of *C. picanus* starting in September using *H. incana* pollen, as well as avocado pollen, as an alternative food, since this Brassicaceae is associated to avocado orchards in the Valparaíso Region and could be used as a refuge in hillside commercial plantations (Bouras and Papadoulis, 2005; Palevsky *et al.*, 2007b).

Mated *A. graminis* females showed a high rate of total consumption, but their eggs were infertile with no descendants over time compared with *C. picanus* that hatched almost 100% of its eggs. Regarding this phenomenon, several authors have pointed out a likeness

Table 3. Total consumption by *Cydnodromus picanus*, *Amblyseius graminis*, and *Galendromus occidentalis* of mobile immature *Oligonychus yothersi* during postembryonic development and longevity of predator mites.

Prey	Phytoseiid state	Phytoseiid species (mean ± SE)		
		<i>C. picanus</i>	<i>A. graminis</i>	<i>G. occidentalis</i>
<i>O. yothersi</i>	Larva	0.00 ± 0.00b	0.00 ± 0.00b	1.73 ± 0.10a
	Protonymph	2.00 ± 0.14b	4.17 ± 0.32a	2.34 ± 0.11b
	Deutonymph	2.63 ± 0.11b	3.93 ± 0.38a	2.50 ± 0.12b
	Unmated adult	59.20 ± 1.66a	39.66 ± 2.02b	34.88 ± 1.34b
	Mated female	142.76 ± 6.34b	298.53 ± 7.44a	90.30 ± 3.49c

Values with different letters in the rows indicate significant differences according to Tukey ($p < 0.05$). SE: standard error.

Table 4. Life table parameters of the *Cydnodromus picanus*, *Amblyseius graminis*, *Galendromus occidentalis* predator mites, and the avocado red mite *Oligonychus yothersi*.

Biological parameters	Species (mean ± SE)			
	<i>C. picanus</i> ¹	<i>A. graminis</i> ¹	<i>G. occidentalis</i> ¹	<i>O. yothersi</i>
R ₀	25.41 ± 1.14b	0.27 ± 0.01d	16.25 ± 0.68c	39.66 ± 1.84a
r _m	0.25 ± 0.00a	-0.06 ± 0.00d	0.19 ± 0.00c	0.22 ± 0.00b
T	12.46 ± 0.16d	20.75 ± 0.31a	14.62 ± 0.25c	16.36 ± 0.19b
λ	1.29 ± 0.00a	0.93 ± 0.00d	1.20 ± 0.00c	1.25 ± 0.00b
n	30	30	30	25

Values with different letters in the rows indicate significant differences according to Tukey ($p < 0.05$).

¹Daily diet = 15 mobile immature *O. yothersi*. SE: standard error. n: number of individuals. R₀: net reproductive rate. r_m: net intrinsic rate of increase. T: mean generation time. λ: finite rate of increase.

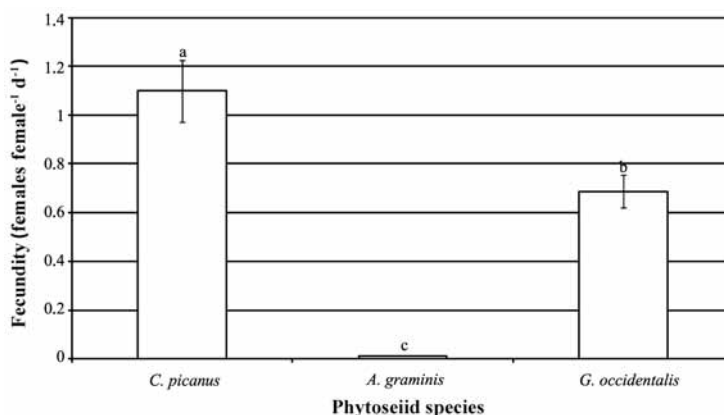
to sequestration or extraction of secondary metabolites from plants, that is, specialist phytophages such as *O. yothersi* would be extracting allelochemicals from the avocado which would then be stored in their bodies as a defense against their predators, thus affecting in distinct ways the three evaluated phytoseiid species (Aregullín and Rodríguez, 2003; Collier *et al.*, 2007; Zhu-Salzman *et al.*, 2008). For this reason, it is necessary to carry out studies to confirm the presence of these toxic substances found in the red mite and predators. On the other hand, phytophages that have a broad range of host plants do not have the capacity to extract these toxic substances (Trigo, 2000; Termonia *et al.*, 2001). Nishida (2002) points out that these substances extracted from the plants are biochemically transformed before being stored in the bodies of lepidoptera. It must also be mentioned that endosymbiotic fungi are present in the plants and influence the tri-trophic interactions (plant-pest-natural enemy), affecting predator development, survival, and reproduction for the production of toxic alkaloids (mycotoxins) (De Sassi *et al.*, 2006).

It must be mentioned that studies evaluating predators based on consumption or female fertility rates do not determine a potential control of the pest and provide incomplete information. High consumption rates do

not imply high female fertility and fecundity, since *A. graminis* showed higher consumption and an oviposition rate similar to *C. picanus* and *G. occidentalis*. However, evaluating fecundity, *C. picanus* had a higher mean of eggs able to develop to the adult stage. It is therefore necessary to determine key biological parameters in ideal conditions to observe the biotic potential of the species of interest.

Establishing life and fecundity tables of predators and prey are fundamental to evaluate the efficiency and potentiality of a natural enemy on a specific pest (Naranjo, 2001; Gabre *et al.*, 2005; Vantornhout *et al.*, 2005; Ozman-Sullivan, 2006; Collier *et al.*, 2007; Reis *et al.*, 2007; Ferrero *et al.*, 2007; Broufas *et al.*, 2007). The above-mentioned information along with consumption registers generate assumptions of potential predator efficiency in the orchard (Chi and Yang, 2003; Kishimoto, 2003; Hosseini *et al.*, 2005; Gotoh *et al.*, 2006). This knowledge is relevant particularly for the assessment of natural enemies that are commercially produced (O'Neil *et al.*, 1998).

In reference to the biological parameters, intrinsic rate of increase (r_m) indicates the capacity of the population to multiply in one generation, relating net reproductive rate (R₀) on generation time (T) (Rabinovich, 1980), implying the potential control of a natural enemy on a specific



Distinct letters indicate significant differences according to Tukey ($p < 0.05$).

Figure 3. Mean female fecundity of *Cydnodromus picanus*, *Amblyseius graminis*, and *Galendromus occidentalis* fed with mobile immature *Oligonychus yothersi*.

pest (Persad y Khan, 2002; Kontodimas *et al.*, 2007). In theory, associating predator intrinsic rate of increase on the prey intrinsic rate of increase, shown by the equation $r_m \text{ predator}/r_m \text{ pest} \geq 1$, will indicate an efficiency potential to regulate the pest population. Other important parameters must also be considered such as longevity, predatory capacity, and early prey detection ability in selecting efficient biological control (Fiaboe *et al.*, 2007). *Cydnodromus picanus* achieved a higher r_m than the red mite, signifying that this population has the capacity to control *O. yothersi* across generations, that is, this species of phytoseiid is an efficient natural enemy of the phytophage mite, and its potential use should be evaluated in the integrated management of avocado mites.

Regarding phytoseiid field releases, all the factors that can influence its effectiveness on a specific phytophage mite must be considered, such as domatia of the host plant (morphological structures of the leaf: depressions, trichomes, cavities between the midrib, and secondary veins that provide refuge for the predator mites generating mutualism) (Matos *et al.*, 2004), chaetotaxia of the predator (length of the dorsoventral setae) (Croft *et al.*, 2004), alternative food availability (Bouras and Papadoulis, 2005), host plant, and leaf area (Collier *et al.*, 2007).

CONCLUSIONS

Given the phytoseiid species under evaluation: *C. picanus*, *A. graminis*, and *G. occidentalis*, it can be concluded that:

C. picanus and *G. occidentalis* complete their postembryonic development and are able to reproduce by feeding on immature avocado red mites in laboratory conditions, both considered as potential predators of

O. yothersi. However, *G. occidentalis* requires prey in the larval stage for its development and without using alternative food.

A. graminis has no descendants when feeding on mobile immature *O. yothersi*. However, its population could be increased in the orchard through a habitat management program since it survives by feeding on avocado var. Hass and *H. incana* pollen as alternative food.

A new predator-prey interaction was established under laboratory conditions (*C. picanus*-*O. yothersi*). Field releases in the spring of *C. picanus* upheld its potentiality as a predator of the avocado red mite in the context of Integrated Pest Management.

RESUMEN

Parámetros de tabla de vida y tasa de consumo de *Cydnodromus picanus* Ragusa, *Amblyseius graminis* Chant y *Galendromus occidentalis* (Nesbitt), sobre la araña roja del palto *Oligonychus yothersi* (McGregor) (Acari: Phytoseiidae, Tetranychidae). En Chile la araña roja del palto *Oligonychus yothersi* (McGregor) es la plaga más importante a nivel foliar en huertos comerciales afectando la fisiología de la hoja, siendo necesaria la búsqueda de nuevos enemigos naturales que interactúen a bajas densidades poblacionales de *O. yothersi*. Se evaluó en condiciones de laboratorio ($27 \pm 1,93$ °C, $87 \pm 3,61\%$ H.R. y un fotoperíodo de 16:8 (L: O)) sobre discos de hojas de palto (*Persea americana* Mill.) var. Hass ($\varnothing = 5$ cm) la potencialidad de ácaros depredadores *Cydnodromus picanus* Ragusa, *Amblyseius graminis* Chant y *Galendromus occidentalis* (Nesbitt), suministrando huevos inmaduros y hembras adultas de *O. yothersi* separadamente, registrando desarrollo

postembrionario, consumo y parámetros de tabla de vida. El desarrollo postembrionario de *C. picanus* fue significativamente menor (5,46 días) en comparación a *A. graminis* (7,33 días) y *G. occidentalis* (8,69 días) al ser alimentados con inmaduros de *O. yotherisi*. Los parámetros de tabla de vida de *C. picanus* fueron tasa neta de reproducción $R_0 = 25,41$, tasa finita de crecimiento $\lambda = 1,29$ y tiempo generacional $T = 12,46$. La tasa intrínseca de crecimiento (r_m) fue significativamente mayor para *C. picanus* ($r_m = 0,25$) frente a *G. occidentalis* ($r_m = 0,19$), mientras que *A. graminis* presentó una $r_m = -0,06$ indicando que su población no tiene descendencia. El r_m registrado por *C. picanus* en condiciones de laboratorio es un indicador del potencial que tiene como depredador sobre *O. yotherisi*, y permite suponer que en condiciones de campo el patrón de reducción poblacional de la plaga podría mantenerse.

Palabras clave: desarrollo postembrionario, depredación, polen, control biológico.

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KINETIC STUDY OF CONVECTIVE DRYING OF BLUEBERRY VARIETY O'NEIL (*Vaccinium corymbosum* L.)

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ABSTRACT

The aim of this research was to study and to model the drying kinetics of the blueberry (*Vaccinium corymbosum* L.) at three temperatures (60, 70 and 80 °C) with an airflow of $2.0 \pm 0.2 \text{ m s}^{-1}$. Modeling of the desorption isotherm was carried out with the GAB (Guggenheim, Anderson and de Boer) equation, showing a good fit to experimental moisture data, giving as a result a monolayer moisture level of $0.084 \text{ g water g}^{-1} \text{ dm}$. Newton, Henderson-Pabis, Page, Modified Page and Logarithmic mathematical models were applied in the study and in the modeling of the drying kinetics of this fruit. Kinetic parameters k of each model showed dependence on temperature, and were evaluated by an Arrhenius-type equation, with an activation energy of between 36.2 and 54.5 kJ mol^{-1} . Logarithmic and Modified Page models gave the best fits for each drying curve, based on the statistical test determination coefficient, sum square error, root mean sum errors and Chi-square. In consequence, both models are excellent tools for estimating the drying time of this product.

Key words: blueberry, GAB, drying, modeling, statistical tests.

INTRODUCTION

The blueberry variety O'Neil belongs to the genus *Vaccinium*, of the family *Ericaceae*, native of North America. It is in the product grouping of berries that includes strawberry, blackberry, raspberry, and others (USHBC, 2007). It was introduced in Chile in 1990 with very good adaptive results, with the result that Chile is now considered as the main producing country of this product in the Southern Hemisphere and the third largest producer worldwide. Of the total production, 85 to 90% is exported, mainly as fresh-cooled, and a small amount is shipped frozen (Chilealimentos, 2007). The production that does not meet whole-berry quality standards is generally converted to clarified and/or concentrated juice, and other technological alternatives are being explored in order to successfully market second quality material, including convection drying with previous pretreatment (osmo-drying, high pressure drying, etc.).

Blueberries are almost spherical, depending on the species and cultivation conditions, and between 0.7 and 1.5 cm in diameter, with a dark blue color. They contain up to 100 very small seeds in a central core (Stückrath

and Petzold, 2007). In Chile, blueberries mature between December and late January, depending on the cultivation zone, and extending over a period of 4 to 5 wk. These berries are a rich source of antioxidant phytonutrients, which are believed to be associated with the antioxidant activity of anthocyanin pigments, flavonoids, and other phenolic compounds containing about three times the total phenolic compounds found in other berries (Skrede *et al.*, 2000). Blueberries have become very popular with consumers because of the research findings that associate their consumption with improvements in human health (Nindo *et al.*, 2007). However, seasonality, market accessibility and costs commonly limit the consumption of fresh fruits and vegetables, so dehydrated products are preferred over fresh products for their convenience, availability and shelf life (Azzouz *et al.*, 2002).

Hot air drying is one of most widely used methods for food preservation. The advantage of dehydrated foods is that decreased moisture content reduces thermodynamic water activity, thus preventing the growth of microorganisms that cause spoilage reactions (Babalís and Belessiotis, 2004). The optimization of dehydration processes in the agro-food industry has led to choosing the technological variables involved in the process itself. Good experimental designs, along with statistical programs, help to obtain a higher yield from the operational and capital points of view. This must be done considering a product of the highest quality, depending on the present variables,

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one of the most important being the time of the drying process. Prior to the study of the drying of any food, it is necessary to recognize and evaluate the moisture sorption isotherms, as these mathematically describe the relation between the water activity and the equilibrium moisture content of the product under study (Vega-Gálvez *et al.*, 2008). The use of mathematical equations is necessary for the simulation of the kinetics of matter transfer (water) that occurs during this unit operation (Vega *et al.*, 2007). There are several empirical equations used to simulate the drying process that are exceptionally practical for the study, the modeling of kinetics and the process optimization, as well as for dryer design (Senadeera *et al.*, 2003). Notably among these equations are those proposed by Newton, Henderson-Pabis, Page and Modified Page, etc. (Doymaz, 2004; Akpinar and Bicer, 2006).

The aim of the present study was to determine and model the drying kinetics and desorption isotherm of the blueberry variety O'Neil, using mathematical equations for both phenomena, and to evaluate the influence of drying air temperature on the kinetic parameters.

MATERIALS AND METHODS

Raw material and proximate analysis

Blueberries of the variety O'Neil (*Vaccinium corymbosum* L.) were cultivated and purchased in the province of Salamanca, Chile. Samples were selected to provide a homogeneous group, based on their date of harvest, color, size, and freshness according to visual analysis. The moisture content was determined according to AOAC methodology N° 934.06 (AOAC, 1990), using a vacuum oven (Gallenkamp, OVL570, Leicester, UK) at 70 °C for 72 h, and an analytical balance (CHYO, Jex120, Kyoto, Japan) with an accuracy of ± 0.0001 g. Crude protein content was determined using the Kjeldahl method with a conversion factor of 6.25. Lipid content was analyzed gravimetrically following Soxhlet extraction. Crude fiber was estimated by acid/alkaline hydrolysis of insoluble residue. Crude ash was estimated by incineration in a muffle furnace at 550 °C. Acidity was determined by the adapted AOAC methodology N° 942.15A (AOAC, 1990), pH was measured using a potentiometer (Extech Instruments, Microcomputer pH-Vision 246072, Waltham, Massachusetts, USA), and sugar content was measured using an Abbe refractometer (ATAGO, 1-T, Tokyo, Japan). All the analyses were made in triplicate and expressed in g 100 g⁻¹ sample.

Isotherm experiments

Desorption isotherms were measured at 60 °C. A known mass of sample (in triplicate) was allowed to come to equilibrium with the atmosphere (relative humidity)

inside a hermetically sealed flask, which contained a glass dish with a saturated salt solution of known water activity. This standard gravimetric method was recommended by The European Cooperative Project COST 90, which deals with the physical properties of foods (Spiess and Wolf, 1983). The weight of the samples was taken every 15 days until reaching constant weight (equilibrium condition). The salts used to obtain a range of water activity of 0.10 to 0.95 included LiCl, KC₂H₃O₂, MgCl₂, K₂CO₃, NaNO₂, KI, NaCl and KNO₂ (Lim *et al.*, 1995). Thymol was added separately in a Petri dish to the recipients containing saturated salt solutions with a relative humidity higher than 75% in order to avoid microbial growth, especially mould (Vega *et al.*, 2007). Once equilibrium was reached, the moisture content of the samples was determined in triplicate.

The relationship between equilibrium moisture content and water activity of blueberries was expressed by the equation proposed by Guggenheim, Anderson and de Boer, commonly termed GAB (Equation [1], Quirijns *et al.*, 2005; Blahovec and Yanniotis, 2008). GAB is commonly used in moisture sorption experiments in different foods, because of its important parameters and physicochemical description, as monolayer moisture (X_m) and the parameters C_o and K_o (Yu *et al.*, 1999; Timmermann *et al.*, 2001). These parameters were obtained by nonlinear regression analysis, using the EXCEL® program of Microsoft® Windows® XP (Redmond, Washington, USA).

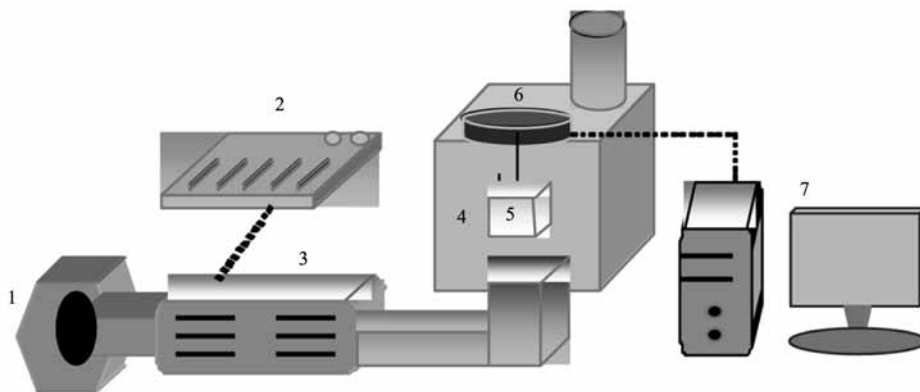
The criteria to evaluate the fit quality of GAB model were the statistical determination coefficient (r^2) and mean percent square error (%E) (Equation [2]) (Vega *et al.*, 2007), where X_{we} is the equilibrium moisture content (g water g⁻¹ dm), a_w is the water activity (dimensionless), X_m monolayer moisture (g water g⁻¹ dm), C_o and K_o are constants of GAB model (dimensionless), X_{ci} is the experimental moisture content (g water g⁻¹ dm), X_{ci} is the calculated moisture content (g water g⁻¹ dm), N is the number of data values and i is the number of terms.

$$X_{we} = \frac{X_m \cdot C_o \cdot K_o}{(1 - K_o \cdot a_w) (1 + (C_o - 1) K_o \cdot a_w)} \quad [1]$$

$$\%E = \frac{100}{N} \cdot \sum_{j=1}^N \left| \frac{X_{ei} - X_{ci}}{X_{ei}} \right| \quad [2]$$

Drying experiments

The drying experiment was carried out using a convective dryer designed and built in the Faculty of Engineering of Universidad de La Serena (Figure 1), La Serena. Three temperatures were used in the study of the drying kinetics (60, 70 and 80 °C). Drying air velocity



1 Ventilator. 2 Control panel. 3 Air heating section. 4 Oven. 5 Sample. 6 Digital balance and interface system. 7 PC.

Figure 1. Schematic diagram of drying equipment.

was held constant at $2.0 \pm 0.2 \text{ m s}^{-1}$ and measured with an omnidirectional anemometer (Extech Instrument Inc., 451112, Waltham, Massachusetts, USA). The inlet relative humidity was $62.0 \pm 5.2\%$, measured by an ambient digital hygro-thermometer (Extech Instrument Inc., 445703, Waltham, Massachusetts, USA). All the drying experiments were carried out in triplicate, using a sample mass of $100.0 \pm 2.4 \text{ g}$ and the charge density was $8.5 \pm 0.3 \text{ kg m}^{-2}$. Prior to the drying of the samples, they were pretreated with a Pectinex® solution, concentration 0.8% (Novo Nordisk Ferment Ltd., Flawil, Switzerland). The mass was measured on an analytical balance (Ohaus, SP402, New Jersey, USA) with an accuracy of $\pm 0.01 \text{ g}$ at defined time intervals, connected by a interface system (Ohaus, RS232, Pine Brook, New Jersey, USA) to a PC, which recorded and stored the data. The experiments were finished at the point of reaching constant weight (equilibrium condition). The dried samples were packaged in polypropylene bags.

Modeling of drying kinetics

Drying kinetic was modeled by means of five empirical equations widely used in most organic and biological materials. These equations use a relationship termed moisture ratio (MR) as a dependant variable (Equation [3]), relating the gradient of the sample moisture in real time (X_{wt}) with initial moisture (X_{wo}) and equilibrium moisture (X_{we}) (Babalís and Belessiotis, 2004). These are Newton (Equation [4]), Henderson-Pabis (Equation [5]), Page (Equation [6]), Modified Page (Equation [7]) and Logarithmic (Equation [8]). In this research, the shrinkage and external resistance were assumed as negligible. Where k is kinetic parameters (min^{-1}), n and a are empirical parameters (dimensionless) and t is drying time (min).

$$MR = (X_{wt} - X_{we}) / (X_{wo} - X_{we}) \quad \text{Akpınar and Bicer (2006) [3]}$$

$$MR = \exp(-kt) \quad \text{Vega et al. (2007) [4]}$$

$$MR = n \cdot \exp(-kt) \quad \text{Vega-Gálvez et al. (2008) [5]}$$

$$MR = \exp(-kt^n) \quad \text{Doymaz (2007) [6]}$$

$$MR = \exp(-(kt)^n) \quad \text{Menges and Ertekin (2006) [7]}$$

$$MR = n \cdot \exp(-kt) + a \quad \text{Akpınar and Bicer (2006) [8]}$$

In order to observe any influence of drying temperature on the kinetic parameters k , an Arrhenius-type equation was applied (Equation [9]), from which the activation energy is obtained ($E_a \text{ kJ mol}^{-1}$), which shows sensitivity of the parameter to temperature (Simal *et al.*, 1996; Simal *et al.*, 2005). Activation energy can be determined by the graphic representation between $\ln k$ versus T^{-1} (K).

$$k = k_0 \cdot \exp(-E_a / RT) \quad [9]$$

Statistical analysis of the models

The fit quality of the experimental data to the desorption isotherm and all the models proposed for drying kinetics were evaluated using the determination coefficient (r^2), sum squared errors (SSE, Equation [10]), root mean sum errors (RMSE, Equation [11]) and Chi-square (χ^2 , Equation [12]) statisticals. The values closest to 1.0 for r^2 , and those closest to zero for SSE, RMSE and χ^2 , are commonly considered as optimum criteria to evaluate the fit quality of the models used (Doymaz, 2004; Akpınar and Bicer, 2006; Doymaz, 2007; Vega *et al.*, 2007). The statistical evaluations were made on the predictions of

equilibrium moisture content by the desorption isotherm, as well as on the drying kinetics. Where MR_{ej} is the experimental moisture ratio (dimensionless), MR_{cj} is the calculated moisture ratio (dimensionless), z is the number of constants and j is the number of terms.

$$SSE = \frac{1}{N} \cdot \sum_{j=1}^N (MR_{ej} - MR_{cj})^2 \quad \text{Simal et al. (2005) [10]}$$

$$RMSE = \left[\frac{1}{N} \sum_{j=1}^N (MR_{ej} - MR_{cj})^2 \right]^{1/2} \quad \text{Togrul and Pehlivan (2003) [11]}$$

$$x^2 = \frac{\sum_{j=1}^N (MR_{ej} - MR_{cj})^2}{N - z} \quad \text{Doymaz (2007) [12]}$$

RESULTS AND DISCUSSION

Experimental analysis and desorption isotherm

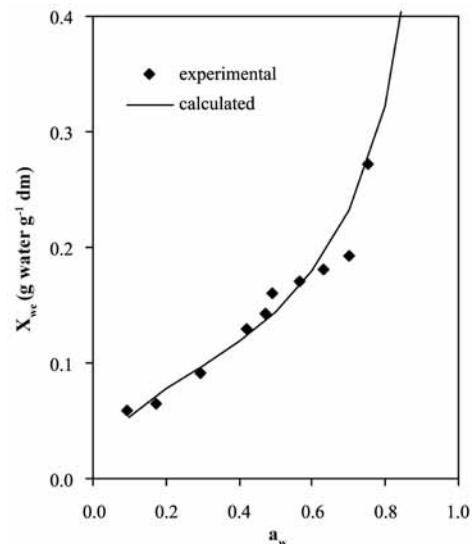
The proximate analysis of blueberry variety O'Neil gave an initial moisture content of 78.13 ± 0.05 g 100 g⁻¹ sample ($a_w = 0.99 \pm 0.01$); crude protein (nitrogen x 6.25) of 0.62 ± 0.12 g 100 g⁻¹ sample; total lipids of 0.40 ± 0.04 g 100 g⁻¹ sample; crude fiber of 6.51 ± 0.27 g 100 g⁻¹ sample; ash of 1.20 ± 0.03 g 100 g⁻¹ sample; non-nitrogen extract (by difference) of 12.90 g 100 g⁻¹ sample; acidity of $2.21 \pm 0.12\%$ (monohydrated citric acid); pH 2.72 ± 0.09 , and soluble solids of 15.01 ± 0.07 °Brix.

Figure 2 shows the experimental data of the equilibrium moisture contents of the blueberry desorption isotherm at 60 °C, where equilibrium moisture content increases as water activity increases from 0.10 to 0.95. The tendency of this isotherm could correspond to that of type II isotherms, according to Van der Waals' classification (Brunauer *et al.*, 1938). Various authors working with other types of foods have observed this behavior (Kiranoudis *et al.*, 1993; Timmermann *et al.*, 2001; Kaymak-Ertekin and Gedik, 2004). The type II isotherm appears when the bonding energy between the water and the primary layer is less than that occurring among water molecules (Lomauro *et al.*, 1985). It was observed the good fit on the experimental data obtained by GAB ($r^2 = 0.97$; %E = 8.64) for a whole range of water activity (Figure 2). The sorption parameters obtained were $X_m = 0.084$ g water g⁻¹ dm, $C_0 = 13.319$, and $K_0 = 0.933$. Similar results of monolayer moisture have been observed in raisins, figs, plums, potatoes, onions and tomatoes with 0.212-0.087 g water g⁻¹ dm for 30-60 °C (Kiranoudis *et al.*, 1993); in red and green peppers with 0.113-0.038 g water g⁻¹ dm for 30-60 °C (Kaymak-Ertekin and Sultanoglu, 2001); in grapes,

apples, potatoes and apricots with 0.220-0.095 g water g⁻¹ dm for 30-60 °C (Kaymak-Ertekin and Gedik, 2004). Monolayer moisture (X_m) is an important parameter since it has a physicochemical behavior that represents the first layer of water molecules, which can thermodynamically interact with other food compounds (Lim *et al.*, 1995; Yu *et al.*, 1999).

Behavior of drying curves

For all drying experiments, an average outlet temperature of drying air of 60.0 ± 4.2 °C was obtained. For this reason, the desorption isotherm modeled by the GAB equation was used to estimate the equilibrium moisture content to each temperature, with $X_{we} = 0.031$ g water g⁻¹ dm for 60 °C; $X_{we} = 0.027$ g water g⁻¹ dm for 70 °C and $X_{we} = 0.026$ g water g⁻¹ dm for 80 °C. In general, for all drying experiments, the equilibrium moisture content was lower than 18%, which gives product stability from the commercial and hygienic points of view (Karathanos and Belessiotis, 1999). Increasing the drying temperature decreased drying time (Figure 3). The shortest time (500 min) was obtained at 80 °C in comparison to drying at 70 and 60 °C, which required times of 800 and 1400 min, respectively. Other research on fruit and vegetable drying present the same tendency and behavior of the drying curves, such as Azzouz *et al.* (2002); Krokida *et al.* (2003); Babalis and Belessiotis (2004); Simal *et al.* (2005); Akpınar and Bicer (2006); Vega *et al.* (2007); and Doymaz (2007).



a_w : water activity (dimensionless). X_{we} : equilibrium moisture content.

Figure 2. Desorption isotherm of blueberries modeled by the GAB equation at 60 °C.

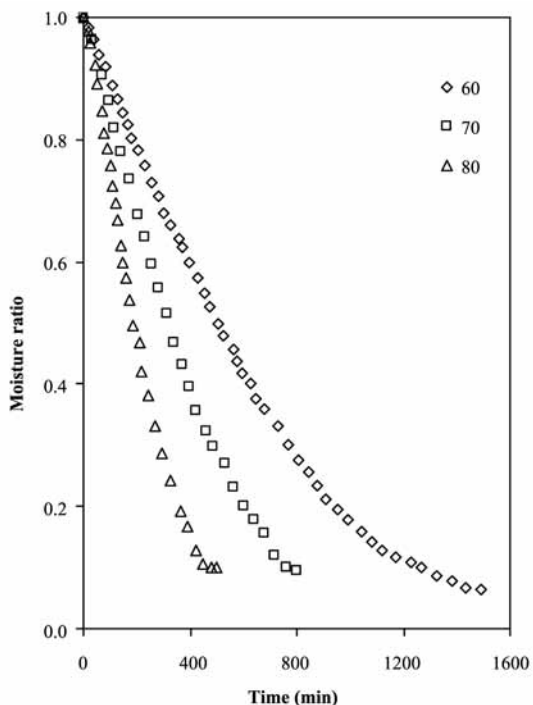


Figure 3. Drying curves for blueberries at different drying temperatures (60, 70, and 80 °C).

Modeling of the drying process

Drying curves (Figure 3) showed a clear exponential tendency and only a falling rate period was observed, which is very common in fruit and vegetable drying processes (Togrul and Pehlivan, 2003; Kingsly *et al.*, 2007). Under these conditions, the use of the five empirical models is suggested, as mentioned before. Table 1 shows the values for the kinetic parameters *k* of the five models for each drying temperature evaluated. A value of *p* < 0.05 for a confidence level of 95% was obtained by ANOVA, using Statgraphics Plus® 5.1 software (Statistical Graphics Corp., Herndon, Virginia, USA), suggesting there are statistically significant differences among these kinetic parameters with respect to temperature. In order to prove the dependence of these parameters on the drying temperature, an Arrhenius-type equation (Figure 4) was applied, showing *r*² ≥ 0.95. The activation energy obtained for each kinetic parameter was 51.05 kJ mol⁻¹ (Newton), 54.45 kJ mol⁻¹ (Henderson-

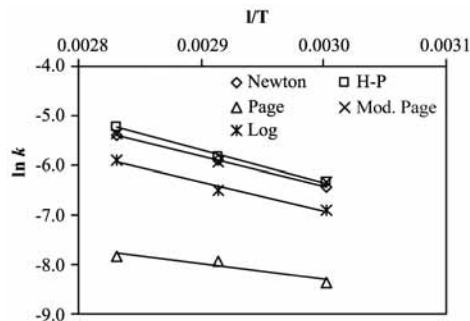


Figure 4. Relationship between the kinetic parameters *k* versus the inverse of temperature (K) modeled by means of five empirical equations: Newton, Henderson-Pabis (H-P), Page, Modified Page (Mod. Page) and Logarithmic (Log).

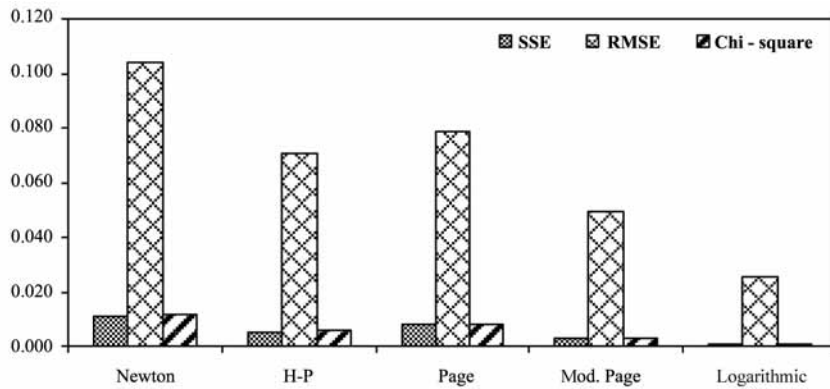
Pabis), 36.20 kJ mol⁻¹ (Page), 46.39 kJ mol⁻¹ (Modified Page), and 48.34 kJ mol⁻¹ (Logarithmic). Several authors have presented very similar activation energy values, including Azzouz *et al.* (2002); Babalis and Belessiotis (2004); Simal *et al.* (2005); Akpinar and Bicer (2006), Doymaz (2007) and Vega *et al.* (2007). Kinetic parameter *k* (min⁻¹) also showed a clear tendency of increasing as the working temperature increased (Table 1). Similar results were obtained by Karathanos and Belessiotis (1999) working with figs, plums and raisins; Togrul and Pehlivan (2003) with apricots; and Akpinar and Bicer (2006) in strawberry drying.

The empirical parameters *n* of Henderson-Pabis, Page, Modified Page and Logarithmic, as well as the empirical parameter *a*, did not show statistically significant differences (*p* value > 0.05), suggesting they probably depend more on the characteristics of the tissue and the drying air flow (Akpinar and Bicer, 2006; Menges and Ertekin, 2006). Azzouz *et al.* (2002), working with grapes, concluded that the parameter *n* was in function of air flow rate and that the parameter *k* of Page depended on the temperature and the initial moisture of the product. Karathanos and Belessiotis (1999), working on skinned and non-skinned fruit dehydration, proposed that parameter *n* increased with the existence of the outer skin depending on its thickness and the kind of product to be dried.

Table 1. Values of kinetic parameters *k* (x10⁻¹ min⁻¹) for each drying curve.

T °C	Newton	Henderson-Pabis	Page	Modified Page	Logarithmic
60	0.0160 ± 0.0002a	0.0177 ± 0.0005a	0.0240 ± 0.0007a	0.0178 ± 0.0001a	0.0101 ± 0.0005a
70	0.0273 ± 0.0003a	0.0300 ± 0.0005a	0.0390 ± 0.0016b	0.0269 ± 0.0002b	0.0152 ± 0.0002b
80	0.0470 ± 0.0005b	0.0557 ± 0.0018b	0.0420 ± 0.0003c	0.0468 ± 0.0011c	0.0273 ± 0.0003c

Data are expressed as mean ± standard deviation of three replications. Values in the same column with the same letter are not statistically different at a confidence level of 95%.



Newton, Henderson-Pabis (H-P); Page, Modified Page (Mod. Page); and Logarithmic (Log); SSE: sum squared errors; RMSE: root mean sum errors.

Figure 5. Graphic representation of the statistic test for each model.

Statistical analyses of models

Logarithmic and Modified Page models provided good fits to the experimental data for a whole drying process (Figure 5). All the models presented high values for the determination coefficient ($r^2 \geq 0.95$) at three temperatures (60, 70 and 80 °C). Furthermore, the lowest SSE, RMSE and χ^2 values were selected as optimal criteria in order to evaluate the fitting quality of the five models proposed. A good fit was observed based on this evaluation, since low SSE (< 0.0027), RMSE (< 0.0495) and χ^2 (< 0.0029) values were obtained by Logarithmic and Modified Page models; followed by Henderson-Pabis, Page and Newton. Other authors have also obtained good results when applying these models in drying kinetics of other food and foodstuff (Krokida *et al.*, 2003; Togrul and Pehlivan,

2003; Doymaz, 2004; Simal *et al.*, 2005; Menges and Ertekin, 2006; Doymaz, 2007).

Figure 6 shows the experimental and calculated values for the drying curves represented by MR vs. time for the two best models found in this investigation (Logarithmic and Modified Page). Both models gave similar results over the whole drying process from the beginning to the end, considering the middle stage of the drying process. The good estimations obtained by these two models represent results that are not usually given by many models, since the middle stage of the drying process is the segment where most of the water is removed from the food. Thus, a good simulation is required for the estimation of the optimum drying time.

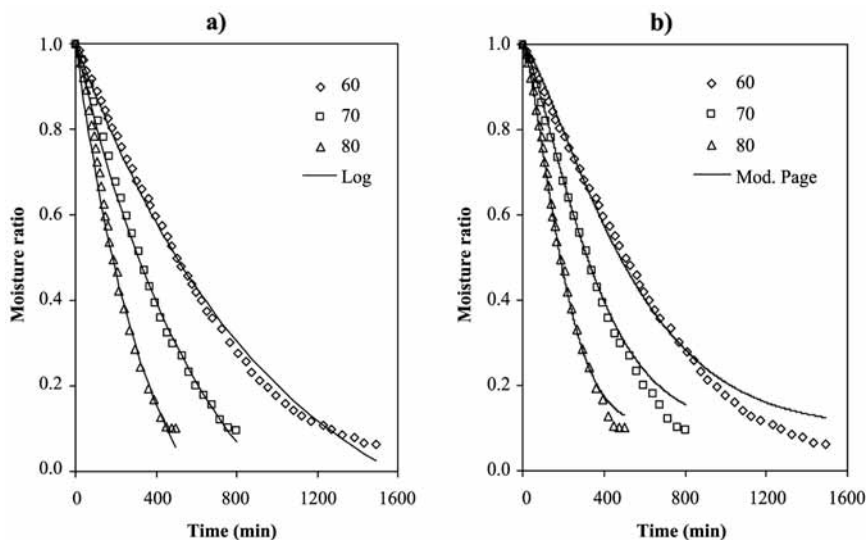


Figure 6. Experimental and calculated drying curves for a) Logarithmic (Log) and b) Modified Page (Mod. Page) at the different working temperatures (°C).

CONCLUSIONS

The results of this study showed that the GAB model provided a good fit to the experimental data of desorption isotherm, with 8.64 %E and $r^2 = 0.97$. Drying of blueberries presents a clear dependence on drying air temperature, showing only a falling rate period, and reaching an average equilibrium moisture close to 0.03 g water g⁻¹ dm. All models used to describe the dehydration kinetics were useful. Nevertheless, Logarithmic and Modified Page models gave the best fit quality to drying experimental data at the three temperatures used, based on the statistical tests used for evaluation. All the kinetic parameters k were dependant on the drying temperature, giving an activation energy of 48.34 and 46.39 kJ mol⁻¹ for Logarithmic and Modified Page, respectively. In consequence, both models are excellent tools for estimating the drying time of this product.

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RESUMEN

Estudio de la cinética del secado convectivo de arándano variedad O'Neil (*Vaccinium corymbosum* L.).

El objetivo de esta investigación fue estudiar y modelar la cinética de secado del arándano (*Vaccinium corymbosum* L.) a tres temperaturas (60, 70 y 80 °C) con un flujo de aire de $2,0 \pm 0,2$ m s⁻¹. El modelado de la isoterma de desorción se llevó a cabo con la ecuación de GAB (Guggenheim, Anderson y de Boer), mostrando un buen ajuste sobre los datos experimentales de humedad, dando como resultado una humedad de la monocapa de 0,084 g de agua g⁻¹ ms. Se aplicaron los modelos matemáticos de Newton, Henderson-Pabis, Page, Page modificado y Logarítmico para el modelado de la cinética de secado de esta fruta. Los parámetros cinéticos k de cada modelo presentaron dependencia con la temperatura, evaluadas por una ecuación de tipo Arrhenius, con una energía de activación entre 36,2-54,5 kJ mol⁻¹. Los modelos Logarítmico y Page modificado obtuvieron el mejor ajuste para cada curva de secado, basado en las pruebas estadísticas como coeficiente de determinación, suma de errores cuadrados, raíz media de los errores cuadrados y Chi-cuadrado. **En consecuencia**, ambos modelos son excelentes herramientas para estimar el tiempo de secado de este producto.

Palabras clave: arándanos, GAB, secado, modelado, pruebas estadísticas.

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LOCATION AND CLASSIFICATION OF MOVING FRUITS IN REAL TIME WITH A SINGLE COLOR CAMERA

José F. Reyes^{1*}, and Luciano E. Chiang²

ABSTRACT

Quality control of fruits to satisfy increasingly competitive food markets requires the implementation of automatic visual servo systems in fruit processing operations to cope with market challenges. A new and fast method for identifying and classifying moving fruits by processing single color images from a static camera in real time was developed and tested. Two algorithms were combined to classify and track moving fruits on image plane using representative color features. The method allows classifying the fruit by color segmentation and estimating its position on the image plane, which provides a reliable algorithm to be implemented in robotic manipulation of fruits. To evaluate the methodology an experimental real time system simulating a conveyor belt and real fruit was used. Testing of the system indicates that with natural lighting conditions and proper calibration of the system a minimum error of 2% in classification of fruits is feasible. The methodology allows for very simple implementation, and although operational results are promising, even higher accuracy may be possible if structured illumination is used.

Key words: fruit classification, color image feature, visual servo control, look-and-move.

INTRODUCTION

When it comes to fresh fruit automatic processing, real time methodologies to accomplish classification tasks are desirable to improve efficiency of fruit processing lines. Visual servo robotic manipulators are able to perform intelligent fruit manipulation based on image plane information in real time if suitable image processing algorithms are provided. Many visual servo control architectures and strategies have been analyzed and classified in the literature (Stavnitzky and Capson, 2000; Xiao and Todo, 2001; Gans *et al.*, 2003), with a mobile camera, either mounted on the effector or in a fixed position to get visual feedback from the workspace. Control strategies can be position-based if control input is defined in terms of absolute position; or image-based if control input is defined according to changes in position of image features on the image plane. The choice of particular servoing architecture relies on aspects such as the inherent geometry of the robotic task, reliability, accuracy,

speed and cost. In many cases the motion of a target, for example an object on a conveyor, is most conveniently expressed in a Cartesian reference frame; therefore most systems dealing with moving objects have used position-based methods (Taylor and Kleeman, 2004; Deng *et al.*, 2005). One of the major applications of visual servo control for robotic manipulators in industry deals with grasping objects from a static or mobile surface, mainly to substitute manual labor in inspection, identification, selection or classification operations (Penman, 2001; Recce *et al.*, 1998). In applications such as processing lines for fruits and vegetable selection, a monocular static camera arrangement is the simplest and most economical implementation, configured to simultaneously image the target fruits and the effector. Color image processing has been used to develop methodologies to assess the maturity levels of fruits. There are discriminator methods based on neural perceptron networks (Leemans *et al.*, 2002) to overcome the fuzzy nature of the class membership identification. Generally, the approach is statistically based and uses a Bayesian type of discriminate analysis (Leemans *et al.*, 2002; Blasco *et al.*, 2003). When these methods make the conversion from red, green and blue (RGB) to hue, saturation and value (HSV) space and the classification functions are multivariable expressions. More computer processing time is required, which is undesirable for real time applications.

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In color segmentation of an image, the objective is to spatially separate regions based on similarities inside each region and differences among distinct regions. The way to undertake the segmentation of color varies from the empirical evaluation of various color spaces (Horiuchi, 2006), to modeling based on physical principles (Gheissari and Bab-Hadiashar, 2003). The primary difference between color segmentation and recognition is that the first utilizes color to separate objects without prior knowledge of the specific surfaces, while the second tries to recognize colors knowing the chromatic characteristics of the surface. Although the two problems are conceptually different; the results of the segmentation can be used in recognition. The concept of color indexing in recognition, implies the use of histograms to index objects in images (Berens *et al.*, 2000), trying to exploit color as a useful aspect for quick detection.

Theoretical considerations

Image to space transformation. Mapping from space to image was done according to proper geometrical modeling of the visual system. Geometric correspondence between space and image is illustrated in Figure 1, along with the camera calibration rig. A point P_0 in space on the calibration rig will be considered, characterized by the coordinate vector $X_c = [x_c \ y_c \ z_c]^T$ in the camera reference frame. The coordinate of P_0 with respect to the calibration rig is therefore $X_0 = [x_0 \ y_0 \ z_0]^T$. Once a spatial view geometry analysis is performed (Reyes and Chiang, 2003), the position of point P_0 in camera coordinate system can be written as:

$$\begin{bmatrix} X_c \\ 1 \end{bmatrix} = \frac{z_c}{d_r} \left\{ K^{-1} \begin{bmatrix} X_p \\ 1 \end{bmatrix} \begin{bmatrix} dx \\ 0 \end{bmatrix} \right\} \quad [1]$$

where: X_c are coordinates of the space point in the camera frame; z_c is the distance from the camera frame origin to the space point along the Z_c axis; d_r accounts for the radial distortion factor; K is the calibration matrix of the camera; X_p are the projected coordinates of the point on the image plane and dx is the tangential distortion vector.

Color feature learning. For a $M \times N$ color image with components (R,G,B) , the chromaticity coordinates (r,g,b) are the normalized components, which are expressed as:

$$\begin{aligned} r &= \frac{R}{R + G + B} \times 225 \\ g &= \frac{G}{R + G + B} \times 225 \\ b &= \frac{B}{R + G + B} \times 225 \end{aligned} \quad [2]$$

where $r + g + b = 255$, and color can be represented in the chromaticity diagram (r,g) (Vertan and Boujemaa, 2000). Given a $M \times N$ color image f characterized by the normalized color components f_r and f_g ; each term of the color distribution histogram matrix H (256×256) can be expressed as:

$$H(r,g) = \sum_{i=1}^{M-1} \sum_{j=1}^{N-1} (f_r(i,j)-r) (f_g(i,j)-g) \quad [3]$$

with $\delta(f-c) = 1$ if $f=c$ or $f=0$ if $f \neq c$. Since the ultimate goal is to recognize an object (effect or object) from the background; a differential histogram H_{OB} between object histogram H_o and background histogram H_B can be evaluated. The H_o histogram is evaluated using Equation [3] from images of objects including its background, while the H_B histogram is obtained by applying Equation [3] to images of the background while it does not contain any object. Each term of the differential histogram H_{OB} can then be written as:

$$H_{OB}(r,g) = \sum_{r=0}^{255} \sum_{g=0}^{255} H_o(r,g) [1 - \text{sing}(H_B(r,g))] \quad [4]$$

where:

$$\text{sing}(H(r,g)) = \begin{cases} 1 & \text{if } (H(r,g)) > 0 \\ 0 & \text{if } (H(r,g)) = 0 \end{cases} \quad [5]$$

The differential color histogram eliminates the color similarities between the object and the background. In order to use the information contained in the differential histogram matrix of Equation [4] for object feature extraction; a subset array $H_M(3 \times n)$, defined as the main components matrix (MCM) will contain the n non-zero values of H_{OB} . Each column of H_M will contain, in the first row, the color frequencies (number of pixels of given color components), and the corresponding color components r and g in the second and third rows respectively. This matrix is organized to have the values of frequencies in descending order in the first row, decreasing to the lowest non-zero value. Consequently, the second and third row contain the r and g components corresponding to each non-zero frequency. Therefore, the MCM for an object may be expressed as:

$$H_M = \begin{bmatrix} nf \\ r \\ g \end{bmatrix} \quad [6]$$

where: nf is a file vector containing the color non-zero frequencies in descending order of magnitude; r and g are file vectors containing the (r,g) pairs corresponding to each color frequency of the vector nf .

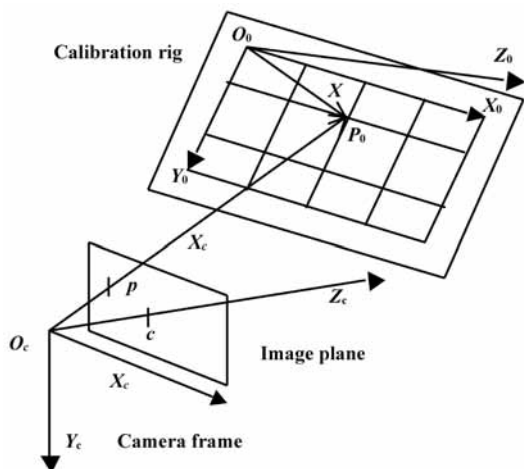


Figure 1. Image and space correspondence.

The objective of this research was to develop a robust and simple methodology that combines color feature extraction algorithms for tracking and classifying fruits as they are moving on a processing line. The main objective was the development and testing of a dual task tracking-classifying algorithm that combines low computational burden with acceptable accuracy to be implemented in commercial fruit processing facilities. The new feature of the methodology is that it employs a single camera, dual-task algorithm suitable for real time applications introducing a new concept in RGB space called a main components matrix (MCM).

MATERIALS AND METHODS

Camera calibration

The images were captured using a charge coupled device (CCD) color webcam pixel view XC75B/465, (Philips, Lisboa, Portugal), with capture card and output image size of 640 x 480 pixels using the standard National Television System Committee (NTSC). The webcam was equipped with a 3.8 mm F 2.0 lens and a viewing angle of 39°. Images were processed with a personal computer (PC) Pentium-S/166 MHz and 64 MB in random access memory (RAM). A geometrical camera calibration procedure was applied to estimate a set of camera intrinsic and extrinsic parameters for mapping between three-dimensional (3-D) space and two-dimensional (2-D) image coordinates (Bouget, 1999). In order to carry out the calibration of the camera, the analytical approach described by Samtaney (1999) was selected as appropriate for the architecture of our simulated system. The procedure employs a minimum of three check board images, 640 x 480 pixels in size, taken at different orientations that are then processed with Matlab software (Bouget, 2001), to obtain intrinsic and

extrinsic calibration parameters. In the present case, 12 calibration images of a check board with 15 x 15, 30 mm squares were used. An additional image calibration for the check board resting on the working plane was added in order to obtain the extrinsic parameters of the task surface, which it is necessary to determine the distance from the camera frame to any point on the working surface.

Object recognition and position estimation

A method for color feature extraction is proposed in order to recognize the class or type of object and determine its position in the image. The method involves two steps. In the first step, the goal is to locate any point belonging to the object. To do this, a matrix $F(M \times N)$ of the same size of the image with zero valued elements is created. Once the image is acquired, color components (r, g) of each pixel are computed and compared against the first, or the first and the second columns of matrix MCM. When equality is found, F is set to one at the corresponding pixel position. The operation can be expressed as follows:

$$F(i,j) = \begin{cases} 1 & \text{if } \begin{cases} f_r(i,j) = H_M(2,1) \cap f_g(i,j) = H_M(3,1) \\ \cup \\ f_r(i,j) = H_M(2,2) \cap f_g(i,j) = H_M(3,2) \end{cases} \\ 0 & \text{if } \begin{cases} f_r(i,j) \neq H_M(2,1) \cap f_g(i,j) \neq H_M(3,1) \\ \cup \\ f_r(i,j) \neq H_M(2,2) \cap f_g(i,j) \neq H_M(3,2) \end{cases} \end{cases} \quad [7]$$

The two highest frequencies of MCM used to recognize the object have to be determined as (r, g) averages of a sample of a group or universe of objects of a certain class in order to have a high probability of finding a pixel of the object image containing any of these two high frequency colors.

Once the entire image is processed with the logical operation of Equation [7], matrix F is searched for the column with maximum non-zero elements (ones) to locate the horizontal position of a pixel inside the object. The sum S_j of all row values for each column of F can be written as:

$$S_j = \sum_{i=1}^{M-1} F(i, j) \quad [8]$$

therefore the horizontal position n_x of the pixel pertaining to the object is equal to the j value such that S_j is maximum in Equation [8]:

$$n_x = j \mid \text{so that } S_j \text{ is maximum} \quad [9]$$

For the vertical positioning in the image, a search along the column n_x is made until the first non-zero value (one) is found. The expression for the n_y position of the point belonging to the object, is then:

$$n_y = i \mid \text{for first value of } F(i, n_x) \neq 0 \quad [10]$$

the second step involves a refinement of the first process, in order to locate the approximate center of area of the image projection of the object. Here more columns of MCM are considered to include the maximum number of chromatic components of the object and therefore the maximum number of image points pertaining to the object. The acceptable number of columns or selective color components from matrix MCM to be used in the recognition process will depend on the degree of accuracy desired with respect to the best estimation feasible. Experimentally it has been shown that at least five columns of MCM are required to obtain a good approximation of the center of area (Reyes, 2002), which means an error of about 2%. The final step employs the pixel position (n_x, n_y) inside the object image, located previously, as the center pixel of a processing window around the object. The window is selected as a $2d \times 2d$ square centered at (n_x, n_y) (Figure 2). To estimate d in pixels, the previous image sampling of the largest size N_T (in pixels) of a certain object class or type, was considered. Estimation of d can then be made as:

$$d = 1.2 \sqrt{\frac{4 N_T}{\pi}} \quad [11]$$

Equation [11] assumes a circular projected object shape, but shape deviation is allowed by increasing the searching region by 20%. The origin of the window positioning (n_{x0}, n_{y0}) is then:

$$\begin{aligned} n_{x0} &= n_x - d \\ n_{y0} &= n_y - d \end{aligned} \quad [12]$$

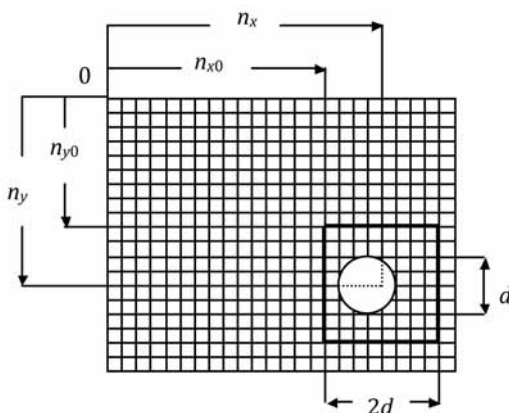


Figure 2. Window for evaluating center of area, d : average diameter of the object; n_{x0}, n_{y0} : coordinates of origin of image window; n_x, n_y : coordinates of center of area of the object in on the image window.

The chromaticity (r, g) of each pixel in the window is compared to the first five columns of the MCM. The respective coordinates (n_j, n_i) of each pixel, with respect to the origin of the window position (upper left corner) are evaluated through the following numerical operation:

$$h_j, n_i = \begin{cases} j, i & \left\{ \begin{array}{l} \text{if } f_r(i, j) = H_M(2, h) \cap f_g(i, j) = H_M(3, h) \\ \text{for } h = 1, 2, \dots, 5 \end{array} \right. \\ 0, 0 & \left\{ \begin{array}{l} \text{if } f_r(i, j) \neq H_M(2, h) \cup f_g(i, j) \neq H_M(3, h) \\ \text{for } h = 1, 2, \dots, 5 \end{array} \right. \end{cases} \quad [13]$$

where: $i, j = 1, 2, 3, \dots, 2d$. If K is the number of pixels where the pair (n_j, n_i) in Equation [13] is non-zero; and n_{jk}, n_{ik} ($k = 1, 2, 3, \dots, K$) are the corresponding coordinates, the approximate position of the center of area of the object projection is calculated as follows:

$$\bar{n}_x = n_{x0} + \frac{\sum_{k=1}^K n_{jk}}{K} \quad [14]$$

$$\bar{n}_y = n_{y0} + \frac{\sum_{k=1}^K n_{ik}}{K}$$

the coordinates obtained in Equation [14] corresponds to the components of the vector $X_p = [x_p, y_p] = [\bar{n}_x, \bar{n}_y]$, which represents the distorted projection of the point on the image plane.

Classification of fruits

This methodology allows discrimination between objects from the same generic class, for example any fruit at some specific state of ripening. In order to implement an algorithm for the identification of objects by color features, a procedure is explored based on the chromatic data contents in the differential histogram defined in Equations [4] and [5]. **The differential histogram targets** the segmentation of the object from the background (separation of the object and background pixels) hereby referred to as the **Background Object Segmentation (BOS)**. First, we group objects according to chromatic similarity (degree of ripening in the case of fruits), by calculating the average (\bar{r}, \bar{g}) of the normalized chromaticity components r and g of each object in a group (fruits with the same degree of ripeness), by means of the expressions:

$$\bar{r} = \frac{\sum_{i=0}^{255} \sum_{j=0}^{255} r(i, j)}{N_s} \left[\text{sing}(H_{OB}(i, j)) \right] \quad [15]$$

$$\bar{g} = \frac{\sum_{i=0}^{255} \sum_{j=0}^{255} r(i,j)}{N_s} \left[\text{sing}(H_{ob}(i, j)) \right]$$

where N_s is the number of pixels obtained from object segmentation. To estimate the average color representative of a given group (similar degree of ripening), the average of averages (\bar{r}, \bar{g}) within the group of n objects is evaluated:

$$\begin{aligned} \bar{r} &= \frac{\sum_{k=1}^n \bar{r}_k}{n} \\ \bar{g} &= \frac{\sum_{k=1}^n \bar{g}_k}{n} \end{aligned} \tag{16}$$

The next step is to evaluate the longest Euclidean distance D_M on the rg plane, between each of the n points (\bar{r}, \bar{g}), corresponding to an object of the group and the point (\bar{r}, \bar{g}), which characterize the group:

$$(D_M)_i = \text{maximum} \left[\sqrt{(\bar{r}_i - \bar{r})^2 + (\bar{g}_i - \bar{g})^2} \right] \tag{17}$$

where $i = 1, 2, \dots, m$ and $j = 1, 2, \dots, n$.

The critical distances $(D_M)_i$ for each group can be employed as a parameter for the classification of any object in one of the m possible groups. To select objects belonging to any of the m groups, we have m components (\bar{r}, \bar{g}) that can be utilized to identify the ownership of each object. The proposed selection criterion involves evaluating the Euclidian distance $(d_o)_i$ in the rg plane from the point $(\bar{r}_o - \bar{g}_o)$ for each object to the reference point $(\bar{r}_i - \bar{g}_i)$ of each one of the m groups:

$$(d_o)_i = \sqrt{(\bar{r}_i - \bar{r}_o)^2 + (\bar{g}_i - \bar{g}_o)^2} \tag{18}$$

where $i = 1, 2, \dots, m$.

To classify any object, the system has to determine which group i satisfies the following relationship:

$$(d_o)_i \leq (D_M)_i \tag{19}$$

On the other hand, to locate and track the object, it is also possible the estimation of the position of its center of area by processing the pixels obtained by means of the Equations [4] and [14].

Fruit recognition

Testing of the procedure was performed using four types of fruits: nectarine, orange, apple and kiwi. The background of the working surface was painted opaque black in order to enhance the discrimination of color features between the environment and fruits. Indirect natural daylight was used from a distant window and filtered by an existing shadowing curtain to avoid reflections caused by direct incident rays. Images of a group of 20 fruits of each type were taken along with an image of the plain background. From each fruit an image window was processed by removing the background pixels and replacing them with pixels of a black color, $(R,G,B) = (0,0,0)$. Files of the modified windows were processed to get average normalized (r,g) histogram defined in Equation [3]. At the same time, a normalized histogram of the background image was also quantified. The differential histogram of each fruit with respect to the background was evaluated by means of Equation [4], followed by the MCM matrix defined in Equation [6]. To locate the center of area of the fruit, a two step computer routine was implemented, where the first step included Equations [7] to [10] and the second, Equations [11] to [14]. These equations yield the two first columns of MCM matrix and the five first columns of MCM respectively.

Experimental classification tests

In order to test the procedure for fruit classification, oranges and apples with distinct color appearance were used. For each type of fruit, 50 fruits with acceptable commercial color were selected and isolated from other group of 50 fruits with deficient color appearance. This procedure was accomplished by using human vision, as is normally done in some manual classification operations. From each of these groups, 11 fruits were randomly grouped to extract color indexes. First the histogram of an image of the isolated background was evaluated, and then the histogram of a window around each fruit. The segmentation of each fruit was finally done by subtracting the average background histogram from the histogram pertaining to each fruit using Equation [4]. For each of the oranges tested, the average of the normalized components (\bar{r}, \bar{g}) was evaluated according to Equation [5]. The following step involved the estimation of the average of averages (\bar{r}, \bar{g}) for each group of similar ripening level, with Equation [16]. Then the maximum permissible distance to this point for each group was determined, by means of the Equation [17]. The averages (\bar{r}, \bar{g}) calculated for oranges and apples, along with the maximum permissible Euclidian distance D_M of Equation [17], were inserted into the algorithm used to test the methodology.

Classification and tracking in real time

Simulated and real tests were performed by means of a multi-thread computer code written in API Windows/C++ that incorporates the analytic formulation of the method described previously. An initial scanning window of 50 x 50 pixels was used along with the dual step MCM algorithm in order to first locate one pixel that belongs to the mobile fruit, and then to estimate its approximate center of area on the image plane. Once position on the image plane is evaluated, the BOS methodology to segment and classify fruit was applied. After object classification was accomplished, the instantaneous center of area was then employed as reference to position and translate a small tracking window of 30 x 30 pixels.

The continuous evaluation of the center of area of a fruit inside the tracking window, along with the image to space transformation, was used to simulate the positioning of a virtual Selective Compliant Assembly Robot Arm (SCARA) manipulator in real time.

RESULTS AND DISCUSSION

Fruit recognition

The averages of five main color components of each type of fruit are shown in Figure 3. Output images in Figure 4 depict the center of area of four fruits (one at a time). In the left image, white crosses indicate the first pixel detected within a fruit. It is important to point out that when there were more than one occurrence (i.e., more than one unit of a type of fruit); the algorithm picked up the object located on the column where more pixels of highest color frequency were found (Equations [7] to [10]). The right frame of Figure 4 illustrates the final output of the method, with white crosses indicating the estimated position of the center of area of every kind of fruit.

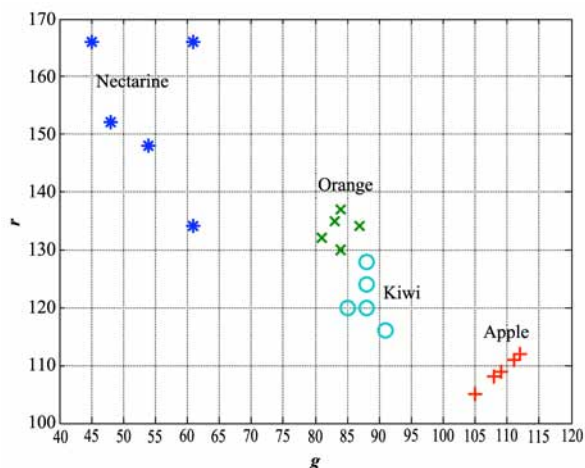


Figure 3. Main color components of fruits, g: green normalized component; r, red: normalized component.

Classification of fruits

In Figure 5 an example of background-object segmentation (BOS) for an orange is illustrated, after applying Equation [4]. **The pixels painted in black are those identified as belonging to the object.** The observed discontinuities are due to noise and color coincidences among the background and the object.

For the purpose of the present analysis, only two regions were identified (Figure 6). The first corresponds to the fruits whose level of ripening is adequate to be accepted (good degree of ripening), while the second region includes those fruits that do not fit the color ripening condition and should be rejected. In this case the objective was testing the method and therefore only two regions were utilized. More generally, it is possible to establish a greater number of regions representing intermediate levels of ripening. **The values (\bar{r} , \bar{g})** for both groups are shown as the center of circles forming each region (Figure 6). The radius of the circle indicates the maximum distance allowed for an object with components (\bar{r} , \bar{g}) to be considered as belonging to the group. Based on this analysis, many groups of fruits can be allocated or classified.

Classification and tracking in real time

A simulated sequence of the method while locating and classifying oranges is shown in Figure 7. Note that the simulated position of the manipulator in the image plane is reproduced virtually using a look-and-move procedure to position the end effector. In the first frame, the border of the initial scanning window is depicted. The second frame shows a random instant captured, while in the third frame the scanning window is still working. In the fourth frame an orange has been classified and tracked with the smaller window.

Numerical results of experimental tests carried out with the method are presented in Table 1. Apples and oranges were visually classified and separated into ripened and unripened. In both cases 50 unripened fruits mixed with 50 ripened fruits were tested using two values of the classification parameter D_M of Equations [17] and [19] previously evaluated using each fruit group. Three repetitions of the test were performed with the same sample. The first value of D_M for each type of fruit (Table 1) is the value given by Equation [17] and the second incorporates a 20% increase in D_M in order to try to improve performance. Outcomes indicated an error fluctuating between 2.0% and 4.6% depending on the value of D_M , and demonstrated that it is possible to get a proper adjustment of the value of D_M for optimum response.

Even though it is possible to adjust the size of the tracking window; the velocity of the center of this window is determined by the sampling rate of the camera. For the

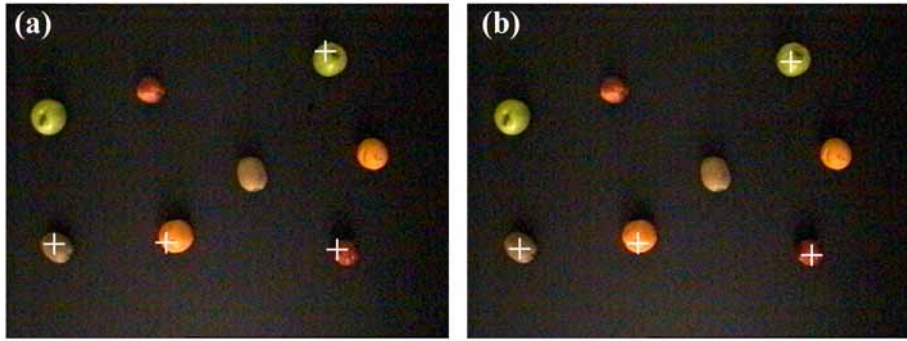


Figure 4. Estimation of center of area for a set of fruits. (a) Determination of a pixel pertaining to the fruit; (b) Determination of the center of area of the fruit.

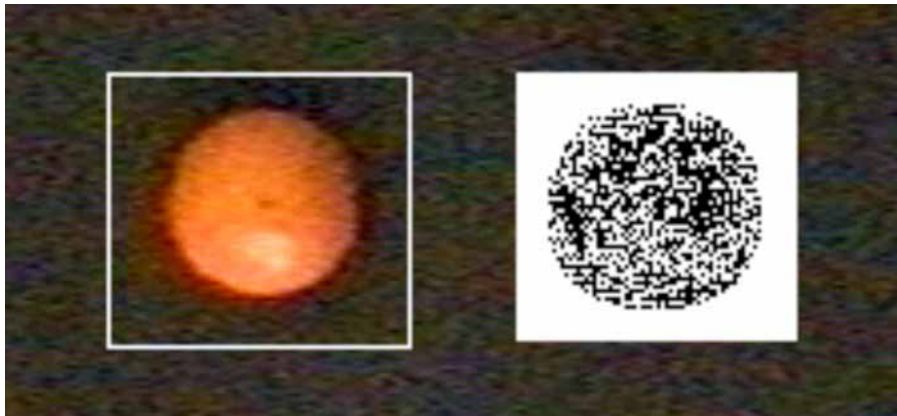


Figure 5. Segmentation of an orange.

trials carried out in this research, typical processing times of each image frame were around 50 to 60 ms, which is less than the speed permitted by the capture rate of our equipment at 66 ms per frame corresponding to 15 images per second. For a square window of 30 pixels, the tracking

operation can be done at a maximum speed of about 450 pixels s^{-1} . This speed corresponds in spatial coordinates to a translation velocity of about 0.5 m s^{-1} , a magnitude that is consistent with the speed of conveying systems used in commercial classification lines for vegetables and fruits.

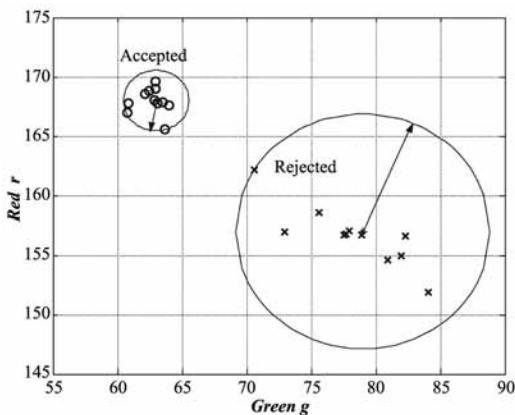


Figure 6. Classification of oranges into groups, g: green normalized component; r: red normalized component. Small circle represents rejected fruits. Large circle represents accepted fruits.

CONCLUSIONS

The methodology presented here uses a single color camera as the sensing device, along with an image to space tracking and classification procedure for fruits. The algorithm proved to be an efficient alternative in classifying and tracking mobile fruits using color feature extraction. The control architecture of a manipulator using the algorithm is beyond the scope of this research, therefore the tracking method was tested assuming a constant moving speed of the fruits. Additional work has to be done in order to evaluate in detail the dynamic behavior of a manipulator employing the methodology developed. Since the proposed methodology was tested under natural lighting conditions, uniform structured lighting may be necessary in order to check operating accuracy of the system. Even though an acceptable

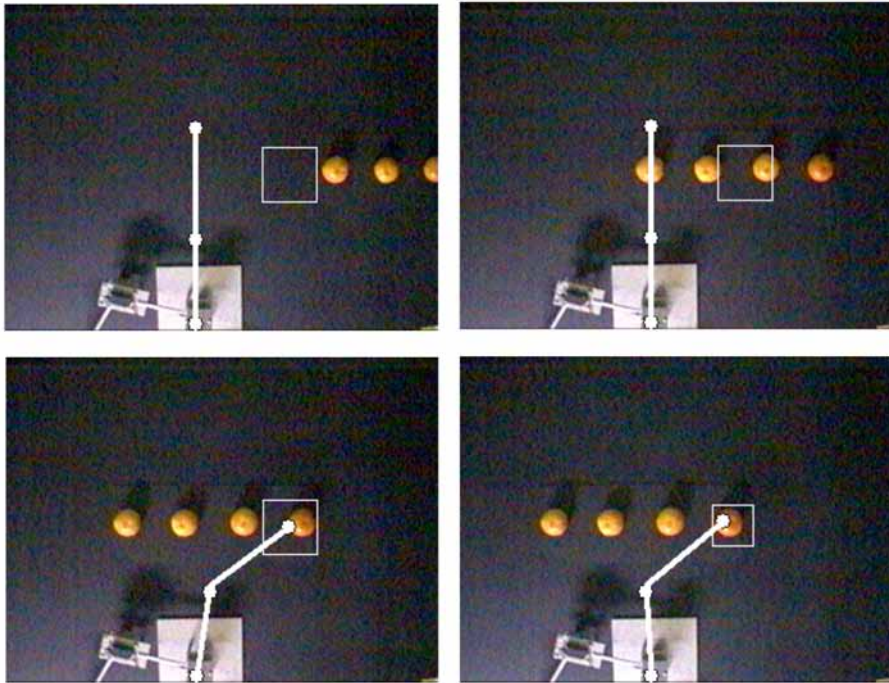


Figure 7. Sequential frames of the simulated methodology captured from the computer screen.

classification error was observed, considerable improvement of the proposed method may be achieved using artificial lighting. In summary, the significant and interesting aspect of this study comprise the deployment of conventional microcomputers and color cameras in devising new automated image-based methodologies to achieve the same objectives as more complex automated architectures.

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RESUMEN

Localización y clasificación de frutas móviles en tiempo real con una cámara individual a color. El control de calidad en frutas y hortalizas para satisfacer mercados cada vez más exigentes, requiere la implementación de sistemas automáticos servo visuales en operaciones de procesamiento de frutas para responder a estos desafíos de mercado. En este trabajo se desarrolló y evaluó un nuevo método para identificar y clasificar frutas en movimiento mediante el procesamiento en tiempo real de imágenes en color capturadas por una cámara individual estática. Se combinaron dos algoritmos para clasificar y rastrear frutas

Table 1. Experimental results for fruit classification tests.

Fruit type	Number of tested fruits	Euclidian distance (D_M)	Number of rejected fruits	Number of failures	Error %
Apples	50	2.0	47.7	2.3	4.6
	50	2.4	48.3	1.7	3.4
Oranges	50	1.5	48.7	1.3	2.6
	50	1.8	49.0	1.0	2.0

D_M , maximum value of the Euclidian distance for a group on the color plane.

en movimiento en el plano de imagen utilizando aspectos representativos de color. El método permite clasificar las frutas en base a segmentación de color y estimar su posición en el plano de imagen, lo cual proporciona un algoritmo confiable para ser implementado en un brazo robótico de manipulación de frutas. Para evaluar la metodología se empleó un sistema experimental simulando una correa transportadora real de movimiento de frutas. La evaluación del sistema indicó que en condiciones de iluminación natural es posible obtener un error mínimo de 2% en la efectividad de clasificación de frutas, con una calibración apropiada del sistema. El método es de implementación sencilla y aunque los resultados experimentales son promisorios se podría obtener una mayor precisión si se emplea luz estructurada.

Palabras clave: clasificación de frutas, aspectos de color, control servo visual, mirar y mover.

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EFFECT OF QUANTITY AND DISTRIBUTION OF RAINFALLS ON *Hordeum murinum* L. GROWTH AND DEVELOPMENT

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ABSTRACT

The growth and development of *Hordeum murinum* L. seeds growing with extreme pluviometric regimes in cool greenhouse conditions were evaluated. Seven treatments according to quantity and distribution of real rainfalls of the semiarid zone of the Metropolitan Region, Chile were applied: rainy-late, normal-late, dry-early, rainy-normal, normal-early and dry-late, plus a reference without water stress, at 2/3 field capacity. The experimental design was randomized complete blocks with five replicate pots. Seeds produced in the last year were sown in pots with disinfected soil leaving the more uniform plants after emergence. Evaluations were made of phytomass production, spearing shoots of roots, the quantity of floral stems and seeds, their total weight and the proportion of seed annex structures, and the viability and germination capacity of seeds. The life cycle of dry years was shortest and with the least dry shoot matter production, the rainy-normal and normal-late years had similar dry root matter production, therefore the most important factor was rainfall distribution. All the reproductive growth values were lower than the reference. There was no seed production in both distributions of dry years and in the normal-early. There were only differences in late distributions, there were no differences among treatments in seed quality. Thus, *H. murinum* uses its resources principally for seed production and late distributions determined seed production.

Key words: pluviometric regime, reproductive efficiency, naturalized poacea.

INTRODUCTION

Annual pasture grasses of the Mediterranean climate of the semi-arid zone of Chile represent the main source of forage for cattle production in areas subjected to water deficits. The magnitude, floral diversity and persistence of the species that make up these ecosystems, as well as the richness of the seed bank in the soil, depend in large measure on the environmental conditions of the location. The phenology and productivity of the species present are regulated by diverse factors, among them are notably the availability of water, which determines the beginning of the life cycle, and temperature, which affects the growth velocity and development of each species. The combination of scarce water resources and high temperatures determines the end of the cycle of annual winter species whose seeds germinate and develop in spring (Castellaro and Squella, 2006). The reproductive stage that precedes the end of the cycle is

vital in the persistence of terophytes, determining the quantity and quality of seeds that are produced (Johnston *et al.*, 2005).

It has been demonstrated that the production and quality of seeds in *Bromus berterianus* Colla (Olivares *et al.*, 2006) and *Erodium moschatum* (L.) L'Her. (Olivares *et al.*, 2004) are strongly determined by precipitation during the reproductive period, given that this determines water availability during the fruiting period. Dry matter (DM) production in this species also depends on the distribution of precipitation and consequently could indirectly affect reproductive parameters through the number of flower stems induced and the provision of carbohydrates assimilated by these (Johnston *et al.*, 2003).

Pluviometric regimes in Mediterranean zones are characterized by high variability in rainfall in both distribution and total levels, because of which native or naturalized species must modify their growth and development, changing, for example, the duration of the periods from emergence to establishment, or the growth rate (Olivares *et al.*, 1998). Acuña (1978) in a pioneering work, simulated pluviometric regimes *in situ* and determined that DM production of annual pasture in a Mediterranean climate increases in those years with

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normal total precipitation, or rainy years, and can also increase or decrease according to the type of distribution of rainfall over the year.

Some seeds of annual species in semi-arid zones only germinate if the level of rainfall is sufficient for the plants to complete this stage and exceed 96 h of hydration. The subsequent growth of these plants will depend on the timing and level of rainfall and the soil water potential (Jara *et al.*, 2006; Castellano and Squella, 2006). Germination and emergence respond to precipitation levels close to or higher than 20 mm in a single rainfall or accumulated in two weeks (Johnston *et al.*, 1998). Rainfall levels in Chile of around 10 mm can be effective for naturalized plants with superficial roots; similar rainfall values were found by Volis (2007) to initiate germination found in *Hordeum spontaneum* Koch. and *Avena sterilis* L. in Israel. On the other hand, higher precipitation levels than these are important for deep-rooting native Chilean species (Gutiérrez, 1993).

As well, it has been established that seed weight is important, in that a greater initial quantity of nutrients generates seedlings of greater vigor, thus assuring a better establishment of the plant and allowing for higher production of seeds (Lorenzetti, 1993). It has been observed that seed weight in ryegrass (*Lolium perenne* L.) is highly variable, depending on climatic conditions under which the seeds develop. As well, climate influences the weight of ovums until anthesis, the moment at which the partition of assimilates is determined (Warringa *et al.*, 1998).

In relation to the availability of water, Kokubun *et al.* (2001) determined that soya plants submitted to water stress produce a lower number of seeds owing to a reduction in photosynthesis; if the deficit occurs during flowering, there is a lower number of assimilates that go to floral structures and consequently there is a smaller number of floral abortions due to damage to the pistils or stamens.

In a study on the effects of pluviometric regimes on the behavior of *B. berterioanus* Colla, one of the most abundant species and of greater forage potential in annual pastures in semi-arid Mediterranean climates, the premise was tested that seed growth and production improved in years with a late distribution of rainfall, whether rainy or normal, and with the dry-early years (Johnston *et al.*, 2005). It can be considered that not only is the quantity of available water important, but also rainfall distribution. Thus, even though there may be less available water in the environment, if this condition is accompanied by other favorable conditions (light and temperature) growth is favored.

Hordeum murinum, one of the most common species in semi-arid Mediterranean ecosystems, shows a high

rate of initial growth, has better adaptive advantages in the context of moderate water deficit and rapidly reaches a seedling size that competes more effectively with the majority of terophytes of the semi-arid Mediterranean range (Olivares *et al.*, 1997).

In accordance with this information, the hypothesis is proposed that the quantity and distribution of precipitation not only influences the growth and development of *H. murinum*, but also has direct effects on the quantity and quality of seeds produced.

Consequently, the general objective of this work was to study the vegetative and reproductive growth of *Hordeum murinum* subjected to different simulated pluviometric regimes in cool greenhouse conditions and determine the production and quality of the seeds of these plants.

MATERIALS AND METHODS

Thirty-five (35) black polyethylene pots were kept under cool greenhouse conditions (without heating), with average temperatures between 15° and 26° and mean relative humidity between 73% and 64%. The pots were 50 cm deep and with a diameter of 13 cm, with 90% of soil previously treated with bromomethane to activate existing seeds and subsequently sieved at 5 mm, plus 10% of poliestirene spheres of the same diameter. Ten seeds of *Hordeum murinum* (synonymous with *Critesion murinum* (A. Löve) were planted in each pot. The seeds had been collected at the Agricultural Experimental Station of the Universidad de Chile, Santiago (33°28' S, 70°51' W). The largest seeds filled with grain were selected, using the most uniform. Seeding was carried out in March in soil with a humidity of 22% dry base weight prior to the beginning of the rains in all of the treatments. Once the simulated rainfalls that corresponded to each treatment began, the number of seedlings per pot was reduced to the five most vigorous.

Given that *H. murinum* and *B. berterioanus* coexist in the natural pasture of the semi-arid Mediterranean climate, the pluviometric regimes were used that had presented positive (+) or negative effects (-) in the production of *B. berterioanus* (Olivares *et al.*, 2006). The amount and timing of the application of precipitation in each treatment were made in accordance to the selected year (Figure 1); the first effective rainfall was considered as the beginning of the period and the following applications were made according to the calendar of real precipitation for the selected year (Gutiérrez, 2003). As well, a reference treatment without water restriction was maintained at 2/3 of field capacity, adding water in quantities determined by the difference in the weight of the pots in relation to the initial content that was equivalent to 50 mm of rain. The soil water content was calibrated on the basis of the data

obtained by Olivares *et al.* (2004) for the same type of soil, from the Cuesta Barriga series (Typic Haploxerolls), with which the soil weight was determined at field capacity (100% of available water). Rainfalls were applied

according to the calendar of each year selected using a serum dispenser that simulates drip irrigation, allowing that the corresponding quantity of water was applied over 24 h.

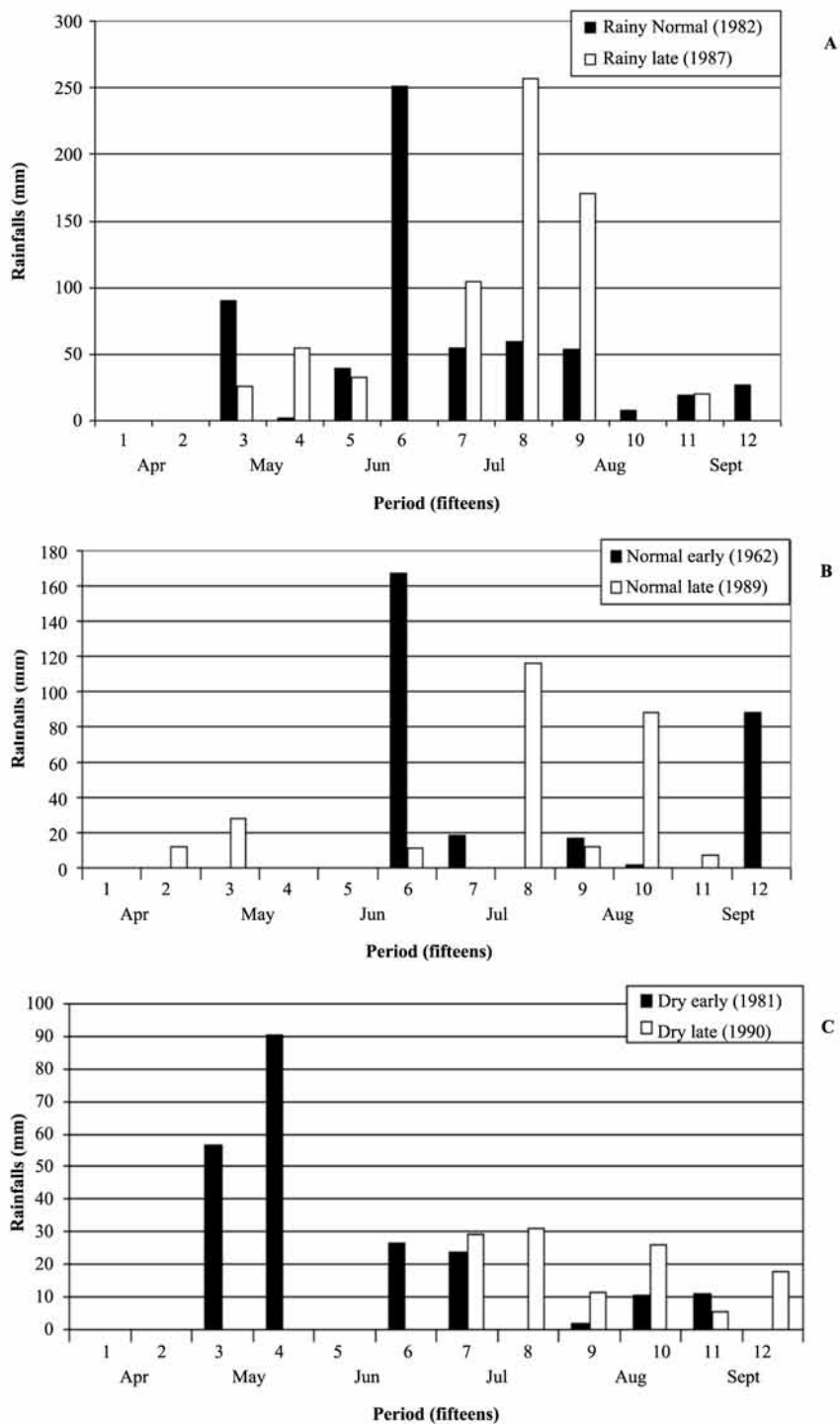


Figure 1. Rainfall distribution (mean of 15 days) of different selected pluviometric regimes: A) rainy years, B) normal years, C) dry years.

The treatments used were: LI-n, rainy year with normal distribution (-) with a total of 628.0 mm (year 1982); LI-ta, rainy year with late distribution (+) with a total of 670.6 mm (year 1987); N-Te, normal year with early distribution (-) with a total of 257.5 mm (year 1962); N-ta normal year normal with late distribution (+) with a total of 281.4 mm (year 1989); S-te, dry year with early distribution (+) with a total of 225.2 mm (year 1981); S-ta, dry year with late distribution (-) with a total of 157.1 mm (year 1990); and the reference without water restriction (control).

Aerial and root phytomass production

This was assessed through the determination of phytomass produced in shoots and roots at the end of the annual cycle and was expressed as the average g DM per pot. To obtain the roots, all the soil from each pot was placed in a mesh that was submerged in water for 24 h. The roots were then separated with a fine rain of water over a screen. The material collected was dried in a forced air oven at 70 °C for 48 h and was finally weighed to obtain the DM of each part. With these values the root/shoot ratio was calculated.

Production of reproductive phytomass

Reproductive phytomass was evaluated at the end of the cycle by counting the number of floral stems once the seeds matured; the quantity of seeds at the harvest date (mature ear) and the total weight of seeds per plant. Seed quality was evaluated by the proportion of palea, lemma and arista (annexes) of seeds per plant and per fruit; by the fullness of seeds by separating well-developed seeds with abundant reserves (full) from those that were not developed (empty); by the weight of 100 seeds as an estimation of size; by viability and germinative capacity. For the latter, germinative tests were conducted with 25 seeds on filter paper and water in a Petri capsule, with eight repetitions per treatment and following the international guidelines for seed tests (ISTA, 2007). To determine the viability of seeds that did not germinate, the seeds were submerged in a solution at 1% of chloride de 2,3,5-triphenyl-tetrazolium for 24 h at 25 °C in darkness, measuring reddening in accordance with the ISTA (2007).

The reproductive index (RI) was calculated for each treatment, which is the ratio between the total weight of disseminules and vegetative phytomass per plant, and the reproductive efficiency index (REI), which is the ratio between the number of disseminules per plant and its phytomass weight (Aronson *et al.*, 1993).

Phenology

The development of the plants was observed and records were made of the main phenological states:

emergence, foliation, tillering, elongation of the stem (a sign of floral induction), flowering, formation and maturing of fruit and senescence of the leaves. Each phenophase was registered when it was present in 50% of the plants in each treatment.

An experimental design with completely random blocks was used, with five replications per treatment. The experimental unit was a pot with five plants. A variance analysis (ANDEVA) was conducted for two ways for each independent variable. When the variation was significant, the means of the treatments were compared using the Newman-Keuls test (Montgomery, 1991). The statistical package Minitab (2000), version 13 was used, with a level of significance of 0.05.

The analysis of the residuals of the analyzed variable (except percentages) satisfied the suppositions of normality and homogeneity of variances, because of which the usual parametric tests to compare treatments were carried out. In the case of percentages, the data was previously transformed to Bliss degrees.

RESULTS AND DISCUSSION

The ANDEVA carried out for each variable studied showed in the majority of the cases that there were major differences between the reference treatment without water restrictions and the treatments with different degrees of water deficit (pluviometric regimes), because of which it was decided to conduct a second analysis excluding only the reference in order to compare the behavior of plants subjected to the studied pluviometric regimes.

Production of aerial and root phytomass

The rainy-late year reached an average value of aerial DM (ADM) per plant (Figure 2) similar to that of the reference treatment (Table 1). The general tendency showed that at lower availability of water the accumulation of ADM decreased; because of which the dry years had the lowest production, followed by the normal-early year. The importance of distribution is notable, given that production in the normal-late year does not differ from that of the rainy-normal year.

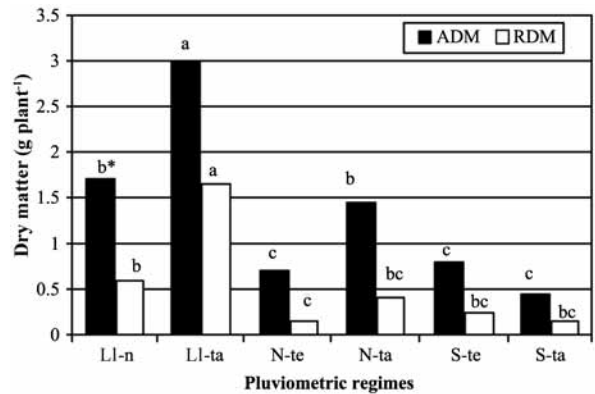
The fact that *H. murinum* can express its growth potential in frequent rainy-late year in the dry interior, given production of ADM was similar to the reference treatment without water restriction, suggests that the pluviometric regime could explain in part the success of this species in this type of year. Other poaceae, such as *B. berterioanus*, which co-exists with *H. murinum* in the pasture, does not reach the same growth potential as it reached without water restriction with any of the prototypes of pluviometric regimes detected in the zone (Olivares *et al.*, 2006).

Table 1. Growth values of *Hordeum murinum* without water stress (reference treatment).

Variable	Values
Dry aerial material, g plant ⁻¹	2.63
Dry root material, g plant ⁻¹	1.35
Root/shoot ratio	0.51
Quantity of floral stems	11.20
Total quantity of seeds	250.70
Fruit weight, mg	1679.00
Seed weight, mg	77.00
Quantity of full seeds per plant	182.20
Quantity of empty seeds per plant	68.49
Weight of annexes, mg	914.00
Percentage of germination	96.00
Percentage of viability	98.00
Estimated size, mm	5.59
Reproductive efficiency index (REI)	0.42
Reproductive index (RI)	62.98

Lower production in dry years could be because of reduced foliar area when there are conditions of water deficit, owing to the fundamental role of water in cellular expansion (González and Páez, 1995). With normal precipitation, early distribution exercises a negative effect, such that there were no differences in the production of ADM in comparison to years with low precipitation with any distribution. From this, it can be deduced that late distribution in rainy years is positive and early distribution in normal years is negative, such that distribution can alter the effect expected based on the total quantity of rainfall. The different timing of rainfalls generated by each distribution results in plant growth occurring at different periods, which can imply more or less thermal and light conditions. This can explain the results obtained and also what is often observed in the field (Castellaro and Squella, 2006).

The DM of the roots (RDM) was less sensitive to the quantity and distribution of precipitation given that higher production was obtained only in the rainy-late year, and the rainy-normal year did not present differences with the normal-late year and dry years with any distribution (Figure 2). The quantity of water and the timing of rains also influenced the root growth of this species, such that while there was a similar behavior among most of the years studied, the most favorable was rainy-late, which produced a quantity of DM equivalent to that produced by the reference without water restriction (Table 1). The normal-late year and the rainy-normal showed similar behaviors despite the major differences in the total amount of rainfall.



* Different letters over columns of the same color indicate significant differences among treatments ($P \leq 0.05$).

Figure 2. Shoot (ADM) and root (RDM) dry matter of *Hordeum murinum* plants according to rainfall levels and distribution. LI-n rainy normal, LI-ta rainy late, N-te normal early, N-ta normal late, S-te dry early, S-ta dry late.

Greater root growth in the rainy-late year corroborates the results obtained with *Bromus pictus* Hook submitted to intense water stress, where the reduction in shoot growth did not significantly alter root growth (Rotundo *et al.*, 2006). Other species also respond to stress by increasing the proportion of assimilates directed to the roots, which allows for diversifying radical growth and thus increase available water for the plant (Huang and Fry, 1998). This explains the similarity of behavior observed in the production of RDM in the different years, with the exception of the rainy-late year. Another work with *Festuca arundinacea* (Huang and Gao, 2000) showed that a moderate stress reduced the length and production of RDM, increasing on the other hand the development of radical hairs with which a greater provision of water is achieved. In studies on the emergence of species of the annual range in the semi-arid Mediterranean zone, it has been demonstrated that *H. murinum* is the most tolerant to a reduction in soil moisture (Olivares *et al.*, 1997). Thus, the results indicate a greater capacity of *H. murinum* to obtain water, or in effect, less sensitivity to water stress.

On the other hand, the fact that a normal-late year has similar production of DM to that of a rainy-normal year, despite the differences in the total quantity of precipitation supports the argument that the distribution of precipitation is very important in the production of phytomass. In this regard, González and Páez (1995) point out that the response to water stress depends on the species and the state of development of the plant at the time when rainfall occurs.

The root/shoot ratio shows similar values among the pluviometric regimes (0.28 to 0.55), with the exception of the N-te, which was significantly higher (1.64); the value of the reference treatment was the same as that the rainy-normal year. This similarity of the ratio in the majority of the conditions studied indicates that normally there would be a greater proportion of resources destined to the shoots than to the roots and that *H. murinum* is capable of adjusting the proportions of phytomass assigned to these structures. The high values of the normal-early year are due to the fact that these plants do not complete their cycle or produce fruits, because of which all of their resources are directed to the growth of shoots. Li *et al.* (2008) points out that climatic variables explain around 50% of the changes in the root/shoot ratio of plants in non-degraded pastures.

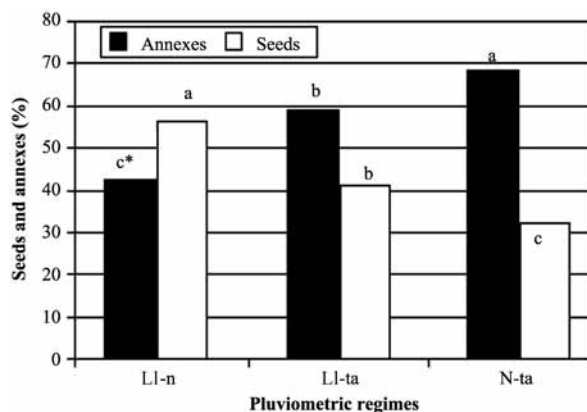
Reproductive biomass production

It should be noted that the dry years, with early and late distribution, and the normal-early year did not reach reproductive development; consequently they are not included in the respective analysis. All the variables on the quantity of measured reproductive growth that completed their reproductive cycle (Table 2) had lower values than those obtained from the reference treatment (Table 1), which would indicate that any water restriction reduces the reproductive structures produced by *H. murinum* L. The behavior was similar in rainy years with both late and normal distribution, while in comparison to the late distributions with different quantities of total precipitation, it was demonstrated that there are differences in all the measured variables, with the normal year being higher.

Kokubun *et al.* (2001) observed that a water deficit in the flowering phase of soya (*Glycine max* (L.) Merr.) reduced the contribution of assimilates to floral structures, determining a higher number of floral abortions and damage to pistils and stamens. Something similar occurred in *H. murinum* in those treatments that did not produce fruits, the similar results obtained in the rainy years indicate that the changes could be attributable, on the one hand, to

factor related to the production and transfer of assimilates, and on the other hand, to the growth and development of seeds, rather than a change in the potential number of seed sites, which coincides with the results of other works (Johnston *et al.*, 2003).

Upon analyzing seed quality, it was observed that the values of the normal-late year were the highest in the number of seeds and weight of annexes (Table 2); in the case of rainy years, although they did not differ among them, they tend to produce more empty seeds and with higher weight of annexes in the late distribution. Considering the relative proportion of annexes and base seeds in relation to total fruit weight (Figure 3) it was demonstrated that the late distributions present higher proportions of annexes, with those from the normal-late year being highest. Nevertheless, in terms of the proportion of seeds, the rainy-normal year had the highest values the normal-late year had the lowest. In the plants without water restriction, there was a high production of seeds (Table 1) with 27% of empty seeds.



* Different letters over columns of the same color indicate significant differences among treatments (P ≤ 0.05).

Figure 3. Annex and seed percentage in relation to total fruit weight from *Hordeum murinum* according to rainfall levels and distribution LI-n rainy normal, LI-ta rainy late, N-ta normal late.

Table 2. Phytomass reproductive components: floral stem (FS), total seed (TS), full and empty seeds number; fruit (seed plus annexes), seed (S) and annex weight of *Hordeum murinum* according to amount and distribution of annual rainfalls.

Treatments	FS	Fruit weight	TS	S	Full seeds	Empty seeds	Peso annexes
	N°	mg	N°	mg	N°	N°	mg
Rainy-normal (LI-n)	1.1b*	58b	16.7b	33b	11.6b	5.5b	25b
Rainy-late (LI-ta)	0.9b	88b	19.7b	37b	11.7b	7.9b	51b
Normal-late (N-ta)	1.9a	210a	45.0a	72a	30.9a	14.1a	142a

* Different letters in the column indicate significant differences among treatments (P ≤ 0.05).

Precipitation and low temperatures during flowering in populations of *Hordeum spontaneum* and *H. vulgare* determined positive correlations between the cross-pollination rate and annual average precipitation and negative correlations between the former and cold temperatures during flowering (Abdel Ghani *et al.*, 2002). On the other hand, according to Egli (2004) water stress during the seed development of oats and other species cuts this period and reduces production. As well, water stress accelerates foliar senescence in *Avena sativa* L. It can then be considered that the normal-late year provides the most favorable conditions for the growth of reproductive structures (fruits), but rainy years favor the movement of assimilates to the seeds.

The pluviometric regime did not affect the capacity for germination, total viability or the estimated seed size (Table 3), and their values did not differ from those presented by the treatment without water restriction. It could be concluded with this that the persistence of the species is assured under unfavorable conditions as argued by Abdel Ghani *et al.* (2002) for *Hordeum spontaneum* and *H. vulgare*. Similar results were obtained by Herrera *et al.* (2008) in *Chloris cucullata* and *C. subdolichostachys* where the percentage of full and dormant seeds changed among years with different distributions of precipitation, but without altering viability.

It has been established that once the number of fruiting sites is fixed, water availability will determine the fullness of seeds, both in terms of production and transportation of new assimilates and the transfer of assimilates from reserve (Johnston *et al.*, 2003). This explains the fact that the reference treatment had the lowest proportion of empty seeds, that the proportion of seeds was higher in the normal rainy year, that the proportion of annexes was higher in late years and that all the seeds produced presented high percentages of germination.

The calculated indicators of reproductive growth (Table 3) show that the pluviometric regimes that complete this phase presented changes in the assignment of resources for the production of disseminules. Thus, the normal-late year was significantly more favorable with

the highest indices, both in reproductive efficiency (REI) and the reproductive index (RI), which indicates that a greater quantity of assimilates are destined to form seeds than in the other treatments, with the exception of the reference, which is also reflected in values of the quantity and weight of the seeds obtained (Table 2).

Works with *H. spontaneum* and *A. sterilis* (Volis, 2007) using gradients of aridity that included quantity and distribution of precipitation showed that increasing aridity reduced the maturation time of the seeds and that there was a negative correlation between the size and number of seeds, a tendency similar to what was shown in the normal-late year.

All of this shows the different strategies that *H. murinum* uses to assign its resources to the diverse components of reproductive growth according to the pluviometric regime that is present, and consequently allows for understanding not only its great productive potential but also its great plasticity, its tolerance to adverse conditions and its great capacity for persistence in the annual range of the semi-arid Mediterranean zone. Something similar was argued by Aronson *et al.* (1993) for some annual desert species, where changes in reproductive biomass in the context of water stress is manifested in both the number and size of seeds, which allows for survival under critical conditions.

Observations of phenological development

H. murinum did not complete its life cycle in the dry years and the normal-early years, while the reference treatment without a water deficit had the longest life cycle and the longest reproductive stage. In the treatments where the cycle was completed a marked separation of the vegetative phenophases was observed. The duration of this stage was similar in all the treatments (Table 4); nevertheless, it can be noted that the rainy-late year had a more prolonged emergence even though its initiation was simultaneous to the other treatments (10 days after seeding). The normal-late year presented a longer tillerage phase than the other treatments and did not have a defined foliation phase.

Table 3. Germination and viability percentage, mean seed size and values of reproductive efficiency: reproductive index (RI) and index of reproductive efficiency (REI) of *Hordeum murinum* according to annual rainfall amount and distribution.

Treatments	Germination	Viability	Size	REI	RI
	%		mm		
Rainy-normal (Ll-n)	95a*	97a	5.6a	6.98b	0.03b
Rainy-late (Ll-ta)	93a	96a	6.1a	4.22b	0.02b
Normal-late (N-ta)	98a	99a	5.8a	21.70a	0.12a

* Different letters in the same column indicate significant differences among treatments ($P \leq 0.05$).

Table 4. Phenological stage duration (days) of *Hordeum murinum*, according to annual rainfall amount and distribution.

Year	Em	Fol	Mac	Fol-til	Stem elong	Flo	Seed	Mat	Senes
Rainy-normal	7	20	7	31	30	37	35	35	46
Rainy-late	25	5	7	25	26	17	12	35	29
Normal-early	15	10	21	14	*	*	*	*	35
Normal-late	10	29	32	0	27	27	23	23	31
Dry-early	17	14	21	10	17	*	*	*	58
Dry-late	11	20	15	17	25	*	*	*	26
Reference	8	23	20	15	44	27	27	54	46

* The totality of the plants did not present the phenological phases, only the beginning of the stages was evidenced.

Em: emergence, Fol: foliation, Til-tillering, Fol-til: foliation of tillerage, Stem elong: Stem elongation, Flo: flowering, Seed: seeding, Mat: maturation, Senes:senescence.

According to Egli (2004) water stress during seed development of oats and other species cuts the cycle and reduces production. It was also observed that water stress in *A. sativa* accelerates foliar senescence.

This species was able to express the total of its growth potential in the tested rainy years and given that its response is very similar to the reference treatment, it could indicate a greater competitive potential in relation to other species of this range, such as *Bromus berterioanus* Colla and *Avena barbata* Pott. which can only present their potential when there are no water restrictions (Olivares *et al.*, 2006; Castellaro and Squella, 2006). Nevertheless, in exceptionally rainy years, these two species have advantages given their potential for growth and development has been higher than what is obtained with pluviometric regimes typical of the area.

The duration of the reproductive stage, considered from stem elongation, was the most affected by the pluviometric regimes and the phenophases showed diverse degrees of superimposition in all the treatments. The year normal-late year had the shortest total duration of the cycle, suggesting that water availability is a determinant of this duration (Table 4).

In general it was observed that without water restriction the species extends its cycle and has a considerably longer seed maturation phase than with the other treatments. The stem elongation phase was also longer and the senescence phase was one of the latest to initiate.

The alterations observed in the duration of the phenophases seems to be directly linked to both the quantity and timing of rainfalls (Figure 1). Thus, the prolongation of the emergence period of the rainy-late year is explained by the fact that between May and June there was little availability of water, which, added to the low winter temperatures, impeded the initial development of the root. On the other hand, maximum precipitation occurring in August shortened foliation in the normal

distribution and prolonged this phase in rainy-late distribution because of a condition of hypoxia or anoxia in the soil, as has been shown in other species (Drew, 1997).

Although the duration of the total life cycles were similar, the rainy years showed different durations of the phenophases in accordance with the distribution of rainfall, particularly in emergence, foliation, flowering, seed formation and senescence. This could be because the water deficit was presented in one case at the beginning and the other at the end of the pluviometric cycle, affecting different development stages in the two treatments. The dry years tested, independent of the distribution, did not allow for reaching the reproductive stage, thus the amount of precipitation will alter the duration of the phases, shortening or extending them according to the quantity of available water.

Although *H. murinum* shortened its phases in the normal-late year in adaptation, it developed a longer vegetative stage in comparison to the reproductive stage, with which it would have a greater probability of generating the photosynthates that assured the higher production of seeds that was observed by Johnston *et al.* (2003) when this species is submitted to stress by cuts. In contrast, a lower quantity of precipitation determined that some phases were not completed, as occurred in the dry-early and dry-late years.

CONCLUSIONS

It can be concluded that:

Hordeum murinum produces a greater quantity of total dry material with more favorable water regimes in terms of the quantity of precipitation and a better distribution.

The distribution of precipitation appears to be the most important determining factor in the production of seeds of this species.

Seed quality, expressed in the capacity to germinate and viability, is not affected by the pluviometric regime of the semi-arid Mediterranean region.

The species assigns a greater part of its reproductive resources to seed production, which gives it more potential for adaptation to distinct water conditions.

In the dry years with early or late distribution of rainfalls and early rainy years there is no possibility of re-seeding for this species.

RESUMEN

Efecto de la cantidad y distribución de las precipitaciones en el crecimiento y desarrollo de *Hordeum murinum* L.

Se evaluó el desarrollo de plantas de *Hordeum murinum* sometidas a regímenes pluviométricos simulados en invernadero frío. En un diseño de bloques completos al azar con cinco repeticiones, se establecieron tratamientos según cantidad y distribución de las precipitaciones reales del secano de la Región Metropolitana, Chile: lluvioso-tardío, normal-tardío, seco-temprano, lluvioso-normal, normal-temprano, seco-tardío y uno de referencia sin restricción hídrica. Se sembraron semillas del año en macetas con suelo desinfectado dejando cinco plántulas uniformes en tamaño y sanidad. Las lluvias simuladas se aplicaron según el calendario del año seleccionado. Se evaluó la producción de fitomasa aérea y radical, la cantidad de tallos florales, de frutos y semillas y su peso total; la proporción de anexos y el llenado de las semillas, su viabilidad y capacidad germinativa. En los años secos el ciclo vital fue más breve y hubo menor producción de materia seca aérea; en los lluvioso-normal y normal-tardío se presentó un crecimiento radical similar, insinuando que es más determinante la distribución de las precipitaciones. Todos los parámetros de crecimiento reproductivo en los regímenes usados fueron inferiores a los de referencia sin restricción hídrica. Los años secos y el normal-temprano no produjeron semillas. La calidad de semillas producidas en el resto de los tratamientos no fue afectada por los regímenes pluviométricos. Se concluye que *H. murinum* destina gran parte de sus recursos a producción de semillas sin alterar su calidad y que las distribuciones tardías son más determinantes para esta producción.

Palabras clave: régimen pluviométrico, eficiencia reproductiva, poacea naturalizada.

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PREDICTION OF THE COMPOSITION OF FRESH PASTURES BY NEAR INFRARED REFLECTANCE OR INTERACTANCE-REFLECTANCE SPECTROSCOPY

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ABSTRACT

Fast and precise analytical tools can contribute to optimize pasture management decisions. This work was carried out to evaluate the potential of one such technique, near infrared spectroscopy (NIRS), to predict the nutritional value of pastures without previous drying of the samples, comparing two forms of collecting the spectra: reflectance, or interactance-reflectance (fiber optic probe). Samples ($n = 107$) from different swards were taken across the humid and temperate regions (Los Ríos and Los Lagos) of southern Chile. Once their spectra were collected, dry matter (DM) and several chemical constituents, such as crude protein (CP), metabolizable energy (ME), neutral (NDF) and acid detergent fiber (ADF), soluble carbohydrates (SC), soluble crude protein (SCP) and neutral detergent insoluble N (NDFIN), were determined as reference data. Calibrations were developed and the best ranked were selected (by cross-validation) according to a lower standard error of cross validation (SE_{CV}) and a higher determination coefficient of cross validation (R^2_{CV}). Calibrations in the reflectance mode, for DM and CP, reached a high R^2_{CV} (0.99 and 0.91, respectively) and a SE_{CV} (6.5 and 18.4 g kg⁻¹). Equations for ADF, SCP and ME were ranked next, with R^2_{CV} of 0.87, 0.84 and 0.82, respectively, and SE_{CV} of 15.88 g kg⁻¹, 15.45 g kg⁻¹ and 0.34 Mj kg⁻¹. Equations for NDF, SC and NDFIN, with R^2_{CV} of 0.78, 0.77 and 0.61, respectively, and SE_{CV} of 35.57, 94.54 and 1.89 g kg⁻¹, respectively, are considered unreliable for prediction purposes. Interactance-reflectance, on the other hand, resulted in poorer equations for all fractions.

Key words: pasture composition; NIR prediction; near infrared reflectance spectroscopy, fresh pastures, fiber optics.

INTRODUCTION

Near infrared reflectance spectroscopy (NIRS) has been widely used as a fast, reliable and multiple method for evaluating the quality of forages, as well as other agricultural products. This technique is based on the absorption properties of the sample in the near infrared (NIR) electromagnetic region, explained by the chemical bonds present in the specimen being scanned, particularly those bonds involving hydrogen (Deaville and Flinn, 2000). The spectrum resulting from the molecular vibration mechanisms can be complicated by a multitude of factors, but with current capabilities, even when the entire spectrum is not understood, it is still possible to extract useful information by employing multivariate calibration methods to construct empirical models that relate relevant spectral variability of a population of samples to its chemical nature (Miller, 2001).

In the case of forages and other materials of agricultural origin, most of the NIRS work has been performed with dry samples, as the high water content in the typical "natural" condition of fresh pasture samples presents some difficulties. Water, quantitatively the most important constituent in fresh forages and other agricultural materials, can be a challenge in the laboratory processing of fresh samples, imposing difficulties in grinding and affecting particle size and shape. Water can also affect the reliability of the detectors in the upper NIR range (Williams, 2001), as it provokes strong absorption signals that overlap and obscure other spectral features and can cause non-linear responses (Reeves, 2000). On the other hand, if samples could be scanned in their fresh, undried state, and their composition or nutritional value predicted within acceptable limits, an approach for fast and reliable predictions in the field could develop, as the industry devises more portable NIR equipment and computers, without sacrificing accuracy. Alternatively, samples arriving at the laboratory could be instantly predicted without the delay of the drying process, which can also alter chemical bonds, affecting the spectrum and, as a result, the perception of some compositional fractions

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(Alomar *et al.*, 1999); Alomar *et al.*, 2003; Deinum and Maassen, 1994. Natural samples, as fresh forage, can be scanned in special rectangular cells, which present a large sample area exposed to NIR radiation in comparison to traditional circular cells. The forage couvette can be inserted in a transport module that allows the sample to be scanned and its NIR reflectance spectrum collected across the long axis of its surface, as the cell is displaced by the mechanism. An alternative to the above is to apply fiber optics technology, which could be interesting when operating under environmental conditions not suitable for sensitive equipment (Osborne *et al.*, 1993). Fiber optics could be attractive for work under field conditions, but the performance of technology on fibrous materials needs to be evaluated. One problem is that optic probes normally have a small scanning area, especially if compared with a large forage cell. Although a higher sampling error could be produced, this could be partially overcome by taking several readings for each sample, assuming that errors occur at random.

The objective of the present study was to assess the potential of predicting the nutritional value of different types of pastures in a fresh, undried condition by near infrared technology, developing calibrations with the spectra taken by reflectance on a large forage cell and by interactance reflectance, by means of a fiber optics probe.

MATERIALS AND METHODS

Pasture sampling

One hundred and seven samples of different types of swards were collected from different paddocks in 13 farms at different locations (39 and 42° S, 72° W) in the temperate, humid Los Ríos and Los Lagos Regions, Southern Chile. Samples were hand clipped at 5 cm from soil, from March 2001 to May 2002, approximately at 1 to 2 wk intervals, covering different seasons, growth stages, predominant species, geographical positions, soil types and other factors that could account for sources of variation in nutritional value and spectral features. The forage obtained represented mixed permanent swards (comprising different proportions of grasses of the genera *Lolium*, *Agrostis*, *Holcus*, *Bromus* and *Dactylis*), but also legumes such as alfalfa (*Medicago sativa* L.) and clovers (mostly *Trifolium pratense* L. and *T. repens* L.) and annual lawn of oats (*Avena sativa* L.) or barley (*Hordeum vulgare* L.). Several broad-leaved species were also present in different proportions.

Spectra collection

Fresh samples, 1 to 2 kg, were taken to the laboratory, cut to 2 to 3 cm with hand shears, thoroughly homogenised

by hand and scanned in a Foss-NIRSystems 6500 scanning monochromator (Silver Springs, Maryland, USA) with accessories (as below) from the same manufacturers, all controlled by a personal computer and software NIRS 3 from Infrasoft International (ISI, Port Matilda, Pennsylvania, USA). Optical data, transformed to microabsorbance units (log 1/R), were stored in suitable files. Spectra were collected either in reflectance, with a large rectangular cell with a quartz window providing 60 cm² of sampling area (Part Number NR-7080, and inserted in a transport module (Part Number NR-6511), or in interactance reflectance with an optic fiber-optic probe (Part Number NR-6775), comprising a double bundle of concentric silica fibers (210 inner illuminators/210 external collectors of diffuse reflected radiation). The probe containing the fiber bundle is protected by an external steel jacket. In the case of the reflectance readings, three scans were taken by rearranging the sample in the cell, averaged and stored. In the case of the interactance reflectance option, samples were packed in opaque polyvinyl chloride (PVC) tubes (25 x 11 cm) with same material caps at both ends and three perforations along main axis, enabling the fiber probe to be tightly introduced to irradiate the sample and collect the readings. In this way, three readings were collected for each sample and stored as described above. As the probe used has the option to adjust the distance between the end of the fiber bundle and the end of the external jacket (path length to the sample), spectra were taken at distances of 0, 0.5, 1, 3 and 5 mm. Accessories for two modes of scanning samples were attached consecutively to the same monochromator as each new batch of new collected samples were scanned. Two events that took place along the experimental period are worth mentioning, as they affected spectral data and eventually the calibrations obtained: the first was a change of the light source (lamp) in November 2001, and the second a routine adjustment in the monochromator in January 2002.

Chemical analysis

After spectra were collected, samples were dried in a forced-air oven at 60 °C for 48 h, ground in a laboratory mill (Thomas Wiley model 4, Arthur Thomas & Co. Philadelphia, Pennsylvania, USA) with a 1 mm screen, and analysed for residual dry matter (DM) using oven at 105 °C for 12 h, crude protein (CP) by Kjeldahl and crude fiber (CF), following AOAC (1996) procedures (method 978.10); soluble crude protein (SCP) after Licitra *et al.* (1996), neutral detergent (with sodium sulfite and alpha amylase) fiber (NDF) after Van Soest *et al.* (1991), acid detergent fiber (ADF) after AOAC (1996) method 973-18, neutral detergent insoluble nitrogen (NDFIN), combining methods for NDF with Kjeldahl, as above; and digestible

organic matter in dry matter (DOMD) by the two-stages *in vitro* digestibility method of Tilley and Terry (1963), modified by incubating (both stages) in an oven at 39 °C in closed flasks. DOMD was in turn used to estimate metabolizable energy (ME) according to a regression on *in vivo* values developed previously in our laboratory (Garrido and Mann, 1981).

Calibrations

Regression equations were adjusted relating spectral data to fractions determined by the reference methods. Calibration models were developed with the software WinISI II from Foss, NIRSystems (Silver Spring, MD), testing different mathematical treatments of the spectra (differentiation order, subtraction gap, smoothing interval), with or without applying Standard Normal Variance (SNV) and Detrend for scatter correction of the spectra. SNV scales each spectrum to have a standard deviation of 1.0 to help reduce particle size effects, and Detrend removes the linear and quadratic curvature of each spectrum (ISI, 1999). The regression method chosen was modified partial least squares.

The same calibration approach was used for spectra collected in reflectance (transport module) and intercanal reflectance (fiber optics). However, while in the first group the full spectra were used (400-2500 nm), in the second the spectra were trimmed, excluding the range of 400-1100 nm, since the detectors of the intercanal reflectance probe are not suitable for that segment.

Cross validation was performed by dividing the set of samples in groups, to adjust the maximum number of terms (to avoid overfitting) and to select the best equations, i.e. those having a lower standard error of cross validation (SECV) and a higher determination coefficient of the cross validation (R^2_{cv}). Four cross validation groups and two passes of elimination of outliers were defined. A critical T value of 2.5 was set for "T outliers", i.e., samples with abnormally high residuals of predicted versus reference values.

RESULTS AND DISCUSSION

Chemical composition

The compositional data for the samples (Table 1) showed a wide variability in composition of analytical data, which confirms the important differences among sampled pastures.

Values for DM in the range of 100 g kg⁻¹ reflect full vegetative growth, typical of mid to late winter. This is also confirmed by unusually high contents of CP, above 330 g kg⁻¹ DM, for this type of plant material. On the other hand, samples from mature swards are also present, with protein contents around or below 100 g kg⁻¹ DM and DM contents in excess of 300 g kg⁻¹. A broad distribution is desirable when a set of samples is selected for the development of NIR calibrations, as a way to have a better representation of the universe to be predicted subsequently in routine analysis.

Spectra

The spectra from samples scanned by reflectance (average of three readings) are presented in Figure 1. The three blocks of parallel spectra that can be seen clearly apart (Figure 1a), are explained by the adjustments on the equipment, as explained earlier. The important base line shift, impressive as it looks, does not necessarily imply that relevant spectral information cannot be extracted by suitable means. If the combined treatments of standard normal variate (SNV) and Detrend are applied, the shifts are no longer apparent (Figure 1b) and some variability appears in different bands. Information can be subsequently refined by mathematical treatments, such as derivatives and smoothing. After applying a first derivative (subtraction) of the spectral data, over a gap of five data points and a smoothing of segments of five data points (Figure 1c), lines tend to lie close together, except in bands where differences are more apparent. This seems to be the case for the segment of 2050 to 2060 nm (Figure 1c, insert), where curves in the lower position (samples 89, 91, 102) had the lowest CP (83.9, 81.2 and 119.0 g

Table 1. Composition of samples obtained by laboratory reference methods.

Fraction	Range	Average	Standard deviation
Dry matter, g kg ⁻¹	92.10 - 359.80	182.20	53.60
Crude protein, g kg ⁻¹ DM	81.20 - 373.20	205.50	65.80
Metabolizable energy, Mj kg ⁻¹ DM	8.82 - 12.47	11.00	0.79
Neutral detergent fiber, g kg ⁻¹ DM	224.30 - 656.60	449.80	77.10
Acid detergent fiber, g kg ⁻¹ DM	162.70 - 375.80	263.00	45.90
Soluble protein, g kg ⁻¹ DM	20.90 - 209.80	91.80	40.00
Neutral detergent insoluble nitrogen, g kg ⁻¹ DM	2.10 - 20.60	7.50	3.50

DM: dry matter.

kg⁻¹ DM respectively) while those in the upper position (17, 29, 33 and 34), had the highest content (355.5, 373.2, 257.2 and 260.6 g kg⁻¹ DM respectively).

Interactance reflectance spectra also showed a base line shift as a result of fixing and regulations on the equipment (Figure 2a). This is no longer apparent after applying a scatter correction treatment (Figure 2b). Besides, changing light aperture produced differences in absorption peaks, with a weaker signal for 5 mm distance (Figure 3), which means that reflected light was more attenuated when it reached collecting fibers.

Calibrations

After testing several mathematical treatments, the best calibrations were selected according to their cross validation parameters. Table 2 shows the statistics of the best calibrations for the different fractions analysed and with the spectra of samples scanned by reflectance and interactance reflectance.

The selected calibrations were obtained with different math treatments. While for reflectance spectra all fractions were best predicted when calibrations were performed

with a first or second order derivative, for interactance reflectance the best equations were developed with the “raw” spectra (with the exception of NDF) and a smoothing for four or five data points. In general, spectra taken by reflectance produced better results than those taken by interactance reflectance for all fractions, with the exception of NDIN, which was similar. The equations obtained in reflectance for DM, CP and ADF showed the highest statistics for certainty, with R²_{cv} of 0.9 or higher, and a SECV lower than a third of the standard deviation (SD) of reference data. This relation between SECV and SD has been proposed as useful for evaluating an equation, which can be considered as reliable for prediction work when SD is more than three times higher than the SECV (Kennedy *et al.*, 1996). Another criterion that can be applied is the ratio between SECV and the average of reference data for a given fraction, and in general the best equations also tended to show values below 0.1 for this relation. This was the case for DM (0.041), CP (0.082) and ADF (0.054), in the reflectance mode. For ME however, although both equations ended with a SECV of less than 5% of the reference data average, they can not be

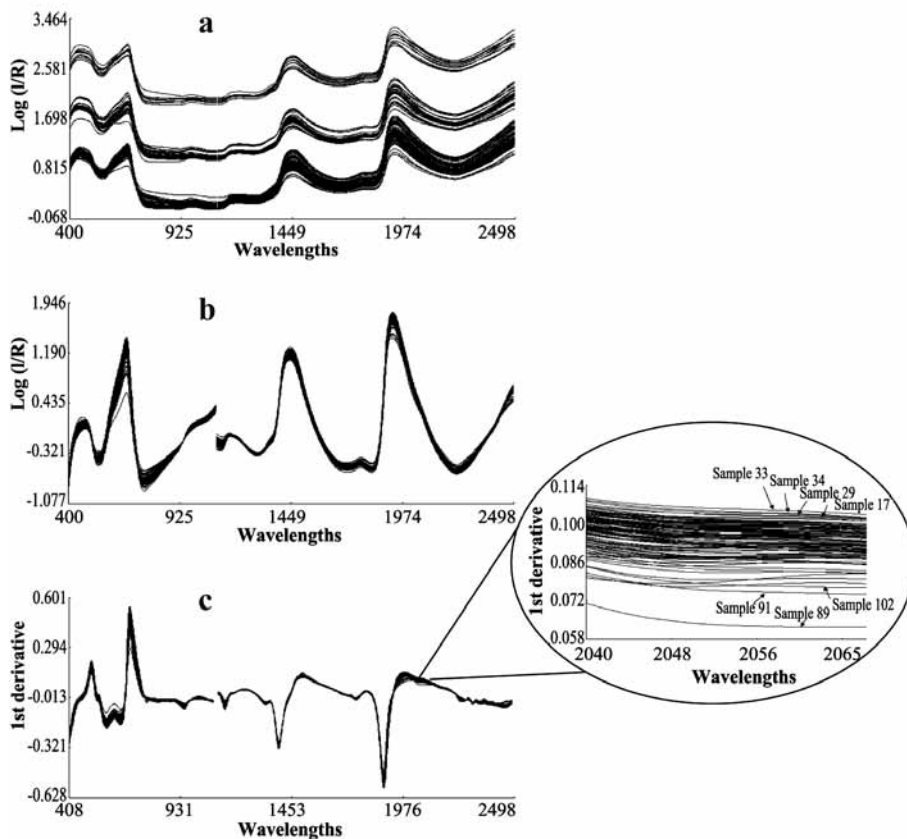


Figure 1. Reflectance spectra of fresh forage from pastures, showing original values (a) or transformed (b) by the scatter correction treatments as standard normal variance (SNV) and Detrend, or the same treatments plus a first derivative (c). Insert depicts a particular segment of the spectra where differences appear among samples of extreme protein content.

Table 2. Statistics of best calibrations with spectra obtained by reflectance or interactance-reflectance.

Fraction	Math treatment*	R ² _{cv}	SECV	SD SECV ⁻¹	SECV Average	Other
Reflectance						
DM	2,5,5,1 SNV+Detrend	0.98	7.50	7.15	0.041	
CP	1,5,5,1 SNV+Detrend	0.93	16.76	3.69	0.082	
ME	2,8,8,1 SNV+Detrend	0.80	0.354	2.23	0.032	
NDF	2,5,5,1 SNV+Detrend	0.80	33.58	2.22	0.075	
ADF	2,10,10,1 None	0.90	13.96	3.20	0.054	
SP	1,10,10,1 SNV+Detrend	0.85	14.44	2.63	0.160	
NDIN	2,5,5,1 None	0.61	1.89	1.59	0.260	
Interactance reflectance (optic fiber)						
DM	0,0,3,1 None	0.84	21.29	2.51	0.120	3 mm path
CP	0,0,2,1 SNV + Detrend	0.75	32.24	2.00	0.160	1 mm path
ME	0,0,4,1 None	0.63	0.47	1.65	0.042	5 mm path
NDF	1,4,4,1 None	0.63	43.87	1.65	0.097	3 mm path
ADF	0,0,5,1 None	0.66	25.53	1.71	0.096	3 mm path
SP	0,0,4,1 None	0.77	18.81	2.11	0.200	1 mm path
NDIN	0,0,5,1 None	0.66	1.76	1.71	0.240	1 mm path

DM: dry matter. CP: crude protein. ME: Metabolizable energy. NDF: neutral detergent fiber. ADF: acid detergent fiber. SP: soluble protein. NDIN: neutral detergent insoluble nitrogen. R²_{cv}: Coefficient of determination of cross validation. SECV: standard error of cross validation. SD SECV⁻¹: ratio of standard deviation of reference data (calibration set) to standard error of cross validation, ratio of standard error of cross validation to average of reference data (see text for details).

* Math treatment: Derivative order (first number), subtraction gap in data points (second number), first smooth segment in data points (third number) and segment for a second smooth segment in data points (fourth number). SNV: standard normal variate (see text for details).

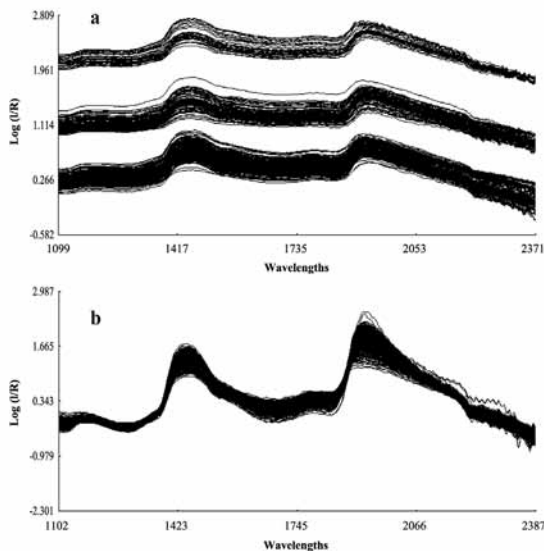


Figure 2. Interactance-reflectance (optic fiber with light path of 3 mm) “raw” spectra of fresh forage from pastures, showing band shifts explained by changes in the equipment (a) and after applying a scatter correction treatment as standard normal variance (SNV), plus Detrend and a first smoothing of five data points (b).

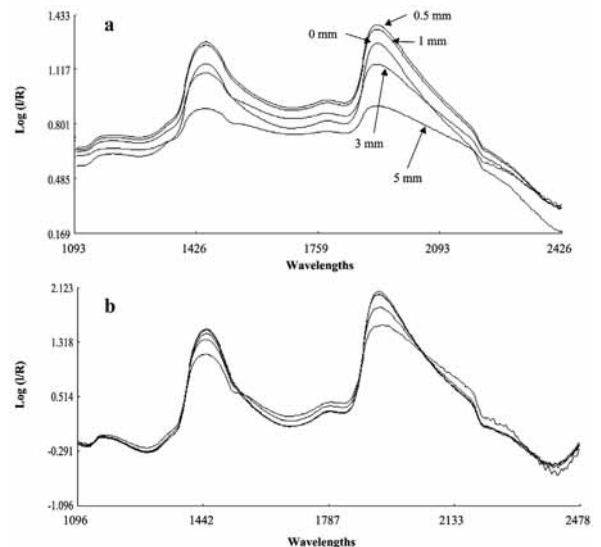


Figure 3. Mean interactance-reflectance (optic fiber) “raw” spectra of fresh forage from pastures. 3a) each curve represents mean spectra for all samples scanned with a given light aperture or pathlength from 0 to 5 mm (arrows); 3b) depicts the same curves, after applying scatter correction treatments as standard normal variance (SNV), and Detrend and a math treatment of smoothing segments of five data points.

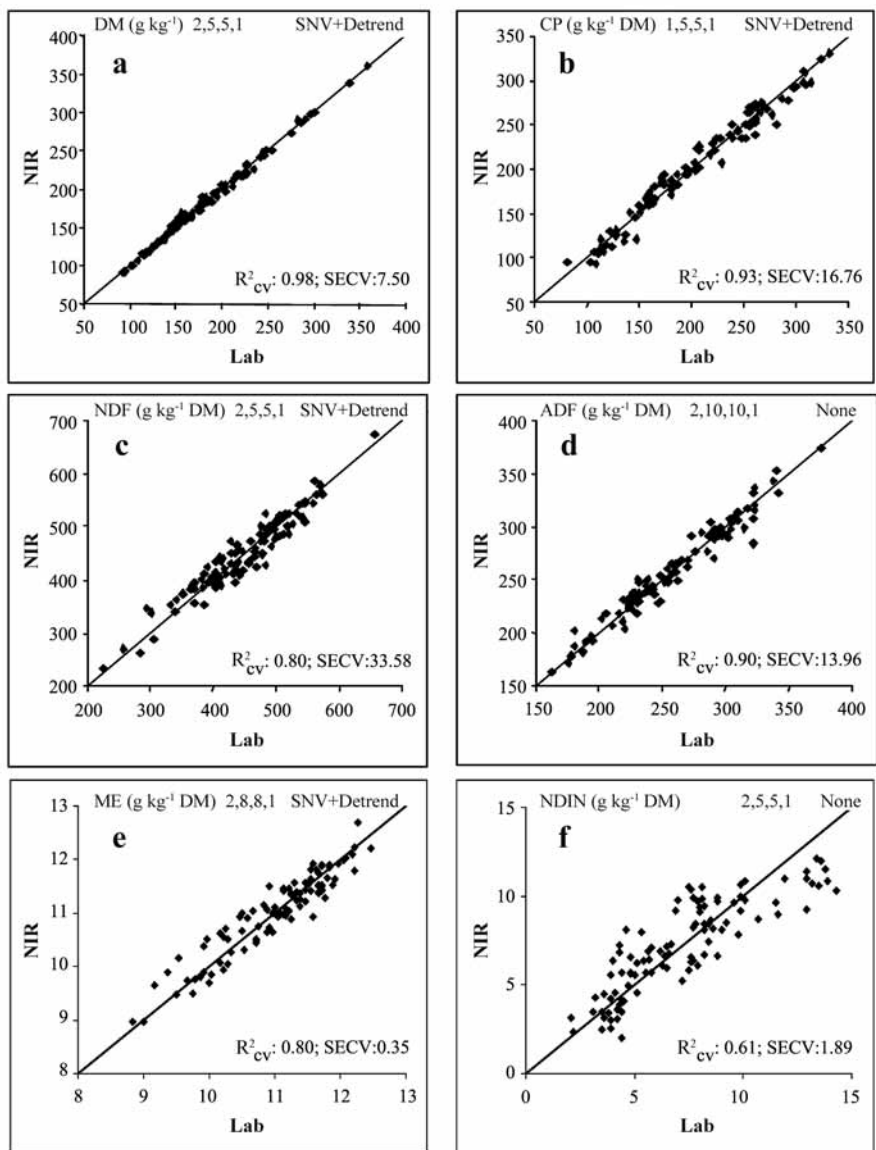


Figure 4. Near infrared reflectance (NIR) predicted versus reference (lab) data for different fractions of fresh forage from pastures. Each graph represents the best calibration obtained for DM: dry matter (a); CP: crude protein (b); NDF: neutral detergent fiber (c); ADF: acid detergent fiber (d); ME: metabolizable energy (e) and NDIN: insoluble nitrogen in neutral detergent (f).

considered as dependable because the error represented an important proportion of the variability of data. This was also the case for the equations for NDF.

The relation between NIR prediction and composition obtained by reference methods was presented for the analysed fractions (Figures 4 and 5). For each fraction, a math treatment was included. For instance, for DM (Figure 4a) a 2,5,5,1 was applied to obtain the best equation, meaning that a second derivative or subtraction over five data points, a first smoothing over a segment of five data points and a second smoothing over one data

point, plus SNV and Detrend, were employed.

It was confirmed that the best equations were those for DM, CP and ADF, in the reflectance mode, as their respective scatter plots depict data points closer to the diagonal equal response line. Although according to these results, NDF and the estimated ME values cannot be confidently predicted by NIRS, a strong relationship can nonetheless be seen in the configuration of the data. This reinforces the idea that spectra can recover signals from chemical bonds that in some way are related to empirical entities, such as those mentioned above.

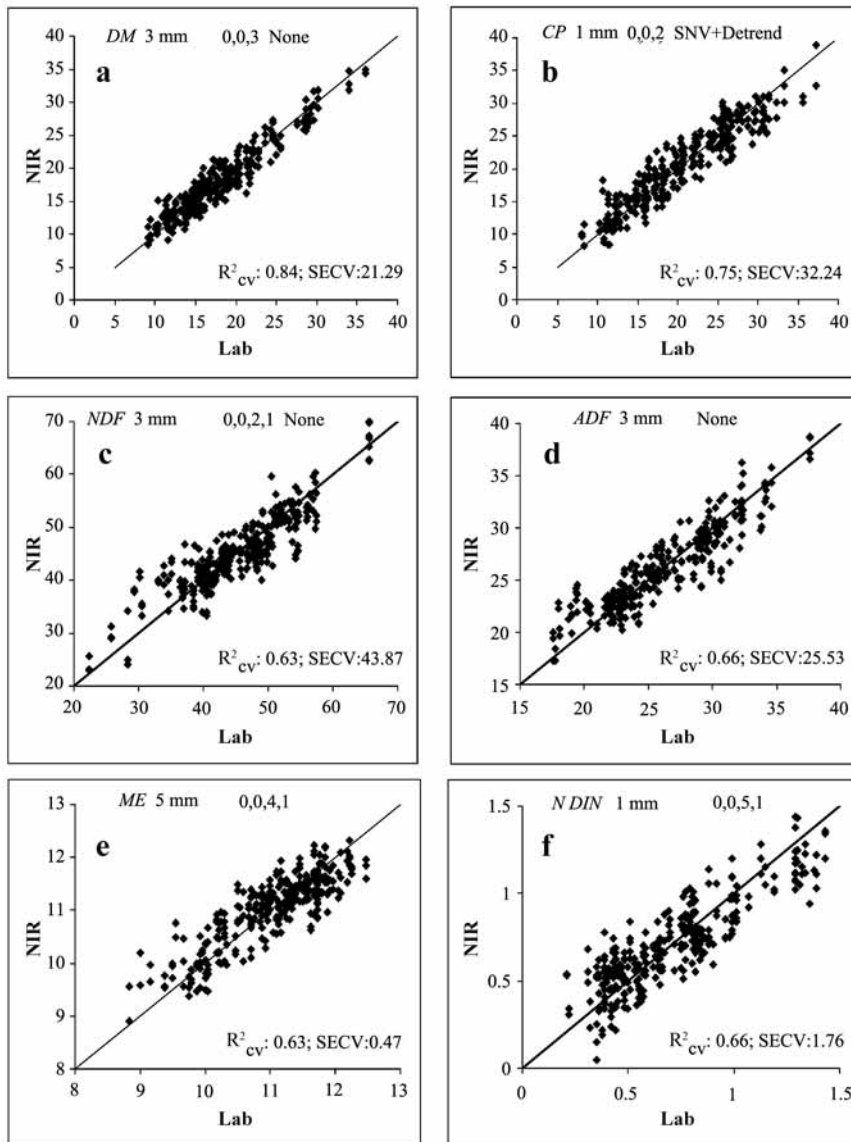


Figure 5. Near infrared reflectance (NIR) (interactance-reflectance) predicted versus reference (lab) data for different fractions of fresh forage from pastures. Each graph represents the best calibration obtained across different math treatments and light paths, for DM: dry matter (a); CP: crude protein (b); NDF: neutral detergent fiber (c); ADF: acid detergent fiber (d); ME: metabolizable energy (e) and NDIN: insoluble nitrogen in neutral detergent (f).

In line with the data presented in Table 2, which shows better results for samples scanned in reflectance, NIR predicted values (Figure 4) are closer to the equal response line (Figure 5). Although a relationship between NIR and reference data can also be distinguished using equations developed from spectra taken with fiber optics technology, the results are far from acceptable for prediction purposes. A probable explanation for these poor results could be in the surface scanned with the optic fiber probe used in this work, which is much smaller than the area covered by the forage cell employed by the transport module for the reflectance spectra. Subsequent work could be oriented

to establishing if a larger number of readings per sample could improve the predictive ability of interactance reflectance technology.

The usefulness of a NIR prediction depends, on the one hand, on the accuracy of the results with respect to reference data, and on the other, on the level of error we are prepared to accept and how fast we can have the results available to make important management decisions. In the case of pasture management, the change in ME and DM content could be important features in deciding when to harvest for forage conservation, or the removal of animal stock from a given paddock.

CONCLUSION

The results obtained in this work demonstrate that several compositional fractions of forage from different types of swards can be accurately predicted by NIRS on fresh plant material, working in reflectance with a suitable forage cell. Fiber optics technology, on the other hand, shows some potential, but results are not acceptable so far.

ACKNOWLEDGEMENTS

This work was supported by a grant from the Chilean National Fund for Science and Technology (FONDECYT), project 1000432.

RESUMEN

Predicción de la composición de pradera fresca mediante espectroscopía de reflectancia o interactancia-reflectancia en el infrarrojo cercano. Disponer de técnicas bromatológicas rápidas y precisas ayudaría a optimizar decisiones en el manejo de praderas. En este trabajo se evaluó el potencial de una de tales técnicas, la espectroscopía de reflectancia en el infrarrojo cercano (NIRS) para predecir el valor nutricional de praderas al estado fresco y comparar dos formas de colectar los espectros: reflectancia e interactancia-reflectancia (fibra óptica). Se colectaron 107 muestras de praderas en las regiones templado-húmedas del sur de Chile (Los Ríos y Los Lagos). Luego de tomar sus espectros, se analizaron por métodos de referencia para materia seca (DM), proteína bruta (CP), energía metabolizable (ME), fibra detergente neutro (NDF) y ácido (ADF), carbohidratos solubles (SC), proteína bruta soluble (SCP) y N insoluble en detergente neutro (NDFIN). Se desarrollaron calibraciones y se eligieron como mejores ecuaciones aquellas que en una validación cruzada, mostraron un mayor coeficiente de determinación (R^2_{cv}) y un menor error estándar (SE_{cv}). Los mejores resultados se lograron en reflectancia para DM y CP, con R^2_{cv} de 0,99 y 0,91, respectivamente, y SE_{cv} de 6,5 y 18,4 g kg⁻¹, respectivamente. Luego se ubicaron las ecuaciones para ADF, SCP y ME, con valores R^2_{cv} de 0,87; 0,84 y 0,82 y SE_{cv} de 15,88 g kg⁻¹, 15,45 g kg⁻¹ y 0,34 Mj kg⁻¹, respectivamente. Las ecuaciones para NDF, SC y NDFIN, con R^2_{cv} de 0,78; 0,77 y 0,61 y SE_{cv} de 35,57; 94,54 y 1,89 g kg⁻¹, respectivamente; resultaron poco confiables para efectos de predicción. La técnica de interactancia-reflectancia produjo resultados inferiores para todas las fracciones.

Palabras clave: composición de praderas, predicción NIRS, espectroscopía de reflectancia en infrarrojo cercano, praderas frescas, fibra óptica.

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MODELLING SUPPLEMENTATION STRATEGIES FOR BEEF STEER REARING AND FATTENING SYSTEMS IN SOUTHERN CHILE

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ABSTRACT

A mathematical model was developed to analyze beef production systems in Southern Chile. The study considered the identification of the main components of systems under different beef steer management strategies, using pasture with or without supplementation and back grounding cattle on pasture followed by a winter period of confined feeding with pasture silage and concentrates. Validation of model outputs using 200 kg LW Hereford steers against real experimental data showed no significant differences ($P \geq 0.01$) between simulated and observed final weights. In order to analyze the interaction between the stocking rate (SR) and supplementation, three SR of 2, 2.5 and 3 steers ha^{-1} with and without pasture silage supplementation at the rate of 5 kg DMd^{-1} steer^{-1} for the length of the entire period were simulated. Means were compared by the least significant difference (LSD, $P \leq 0.05$). Significant differences were found in terms of final weights, which decreased with increasing SR regardless of the supplementation level, although silage supplementation tended to reduce differences between SR. A second set of simulation runs was carried out to simulate on-farm finishing of the steers through a final phase of confined feeding based on a ration of silage and concentrates. Final weights differed between SR and systems and results showed that the optimum corresponded to 2.5 steers ha^{-1} , since at this SR the largest income corresponded to the smallest mean cost. It is concluded that a stocking rate of 2.5 steers ha^{-1} is feasible if winter supplementation is available, independently of a finishing period in feedlot.

Key words: beef cattle, supplementation, grazing systems, simulation models.

INTRODUCTION

Beef production systems in Southern Chile ($38^{\circ}41' \text{ S}$, $72^{\circ}25' \text{ W}$, 200 m.a.s.l. and 1300 mm rainfall) are based on dual purpose (beef and milk) cattle. The predominant breed is Overo Colorado, crossed with Hereford and occasionally other beef breeds. Traditionally, dual purpose breeds account for nearly 80% of the total cattle population (Catrileo and Goic, 2005). According to official statistics, nearly 50% of the total dairy and beef cattle population is owned by peasant farmers (ODEPA, 2001) that run predominantly cow-calf systems with low levels of productivity and efficiency. The overall stocking density is 0.5 cattle units ha^{-1} devoted to livestock, on farms of less than 50 ha where the main forage resources are naturalized pastures and hay supplementation in

winter (Catrileo and Klee, 2005). Naturalized pastures are generally old perennial ryegrass (*Lolium perenne* L.) white clover (*Trifolium repens* L.) grasslands that are degraded to various extents over time and are mixed with a variety of volunteer grasses. Pasture growth is markedly seasonal, with 60% of the total pasture yield concentrated in spring, with a small second peak in the fall.

Animal are reared, backgrounded and fattened on year-round grazed pastures. Pasture-fed finished steers can be produced if sown, well managed pastures are available, but most calves produced by small farmers are sold for backgrounding and finishing to medium and large farms that have better resources, such as sown pastures and cereal crops, and which are therefore able to finish cattle for the internal and export market. This group is the focus of the current paper.

Regardless of the situation, there is general agreement that well managed perennial ryegrass and white clover pastures are consistent with animal well being and preservation of natural resources and provide cheap feed.

Independently of size, farmers generally use rotational grazing and supplement pasture hay or silage to

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grazing animals during the low forage availability winter period. Under these conditions, conserved forages are a relatively expensive feed, and the interaction between conservation, supplementation and the on-farm stocking rate is hypothesized to have an important influence on the economic outcome of the production system.

The favourable climatic and sanitary conditions of Southern Chile, including the fact that the country is free of foot-and-mouth disease, and the relatively free access of Chile to the major beef importing countries, constitute an important incentive for the intensification of beef production. Nevertheless, weak within-country production chains and links, and the fact that most beef farms are relatively small, conspire against the expedient incorporation of new technologies and a sustained process of intensification.

The many possible combinations of variables, such as farm size, initial weight and age of animals, initial grazing date, grazing management and feed quality are associated with the economic outcome of the system. Nevertheless very few technological options have been experimentally tested, and therefore the present study used simulation to analyze the effect of changes in critical farmer-controlled system variables on the likely physical and economic performance of medium-size farms, and assess the probable impact of introducing new technological options.

The objective of the model was to evaluate the productive and economic response of the animals through different scenarios feasible to find in rearing-finishing beef systems in order to improve the income of the system.

MATERIALS AND METHODS

To develop the simulation model it was necessary to identify the main system components, subsystems and relevant variables and write the computer code, as briefly described below.

Model development

The development of the model followed the system analysis methodology advanced and used by several authors (Hoover and Perry, 1989; Aguilar *et al.*, 2003). The model and databases were adapted to simulate performance of beef cattle grazing temperate pastures in Southern Chile. The model was programmed in Visual Basic NET (Halvorson, 2003).

The model used input and output variables (Aguilar, 1997). Input variables included: number and weight of steers; type, quantity and nutritive value of pasture and its size, quantity and nutritive value of supplement, as well as the timing of supplementation; market conditions,

labor and salaries, available infrastructure and its cost. Output variables were: dry matter (DM) intake, stocking rate, final live weight (LW), LW gain; and average and marginal costs.

State variables included: digestibility of available DM; diet selection represented through a selection index, corrected for availability and digestibility; potential DM intake, crude protein (CP) and metabolizable energy (ME); CP and ME concentration of the consumed diet; energy costs of grazing and maintenance, the final energy balance that results in LW gain or loss. Daily feeding costs, including that of the pasture and eventual supplement, are computed, as well as related labor costs, financial costs due to animals, and the proportional costs due to infrastructure and taxes.

The model comprises four main modules: animals, pasture, management and market conditions. The animal component considers forage intake, nutritional requirements for maintenance, LW gain and grazing, and the energy balance that allows normal animal performance according to physiological conditions. The pasture component uses empirical growth curves determined for pastures and forages commonly used in the region. The respective module reads the average monthly potential pasture DM growth and its digestibility, and these initial values are modified according to the state of the system. The management module allows for rotational grazing, the optional use of supplements, and also represents the use of infrastructure and labor characteristic of the region. Market conditions include estimates of fixed costs like labor, infrastructure and equipment maintenance, and variable costs, such as feed and animal health.

Biological components

Animals. The endpoint of the finishing process is essentially conditioned by the animal's weight, but the age at slaughter influences meat quality and needs to be considered as well. Therefore, both variables determine the sale of the animals during simulation runs, and the user can specify age and/or minimum sale weight.

Estimation of the requirements for ME and CP were based on equations proposed by the Agricultural and Food Research Council (AFRC, 1993) and the Commonwealth Scientific and Industrial Research Organization (CSIRO, 1990). The amount of these nutrients consumed is based on the intake of forage and/or supplement, as advanced by Aguilar *et al.* (2003) and CSIRO (1990), whereas the interaction between forage consumption and concentrate intake was based on Aguilar (1997). Voluntary pasture intake and cattle mortality were treated as stochastic variables with user-specified mean and standard deviation.

The model estimates daily intake of forage and supplement under grazing conditions. Initially, a potential DM intake (PDMI) is calculated reflecting the characteristics of the animal under non-limiting conditions. To this end, the model uses a reference animal weight, the current weight of the animal, and its relationship to mature weight (CSIRO, 1990). An algorithm (Aguilar *et al.*, 2003) that computes substitution, supplementary, or additive effects between pasture and supplement intake is used to correct voluntary forage DM intake (VFDMI). The latter is further corrected by a digestibility factor (FCG) and an availability factor (FCD) as follows:

$$\text{VFDMI} = \text{PDMI} \times \text{FCG} \times \text{FCD} \quad [1]$$

where FCG is estimated based on the forage availability, and FCD is calculated from a linear equation adjusted for digestibility values between 40 and 80%. The DM digestibility of diet is estimated from a selection index that corrects the average digestibility of offered forage (Aguilar *et al.*, 2003).

Pastures. Monthly potential growth rates of the pastures, and their variation in digestibility and CP, were used as an input to quantify the potential pasture's DM and nutrient availability. Forage digestibility was corrected for selective grazing, which in turn depended upon forage availability and stocking rate, as discussed by Aguilar *et al.* (2006). For finishing the animals, the silage requirement was simulated in the model assuming *ad libitum* intake and typical nutritive values observed at the farm level.

Economic analyses

Quantification of the economic performance of alternative systems is essential for comparing production alternatives. The simulation model therefore considers fixed and variable costs of production. Fixed costs include labor, infrastructure and machinery maintenance, and general administrative costs. Variable costs are those related to feeding and animal health. The opportunity cost of land was considered as a third component, quantified as the cost of producing forage, and that of animals. Thus, the user has access to two economic analyses, a total and an operational cost, depending upon the inclusion of opportunity costs or not.

Statistical analysis

Initial and final conditions for the experimental and simulated results were compared and the final weights were compared to validate the model using a z test, assuming homogeneity of variances and with a sample size of 40 animals.

In order to analyze the interaction of the stocking rate and supplementation, three stocking rates were simulated with and without grass silage supplementation at the same rate. Each scenario was replicated 20 times. Means of the output variables were compared by the least significant difference (LSD, $P \leq 0.05$).

RESULTS AND DISCUSSION

Model verification

Adequately documented experimental results useful for a thorough validation of model outputs are very scarce. Rojas and Romero (1994) carried out a grazing experiment replicated in each of five consecutive years, in the Central Valley of La Araucanía Region, Chile, using 7-8 mo of age, 200 kg LW, weaned Hereford steers introduced to the experimental pastures on April of each year and fattened until December, using an intensive system of rotational grazing. Simulation runs were carried out for a similar length of time as the experiment, and with the same amount of silage supplementation. No significant differences ($P \geq 0.01$) were found between simulated and observed final weights. Similarly, there were no significant differences ($P \geq 0.05$) in the amount of silage produced in each of the 5 years (Table 1).

Experimentation

To study changes in economic and physical performance of the system in response to management practices, a number of scenarios, including changes in the stocking rate, levels of supplementation during the winter period, and a final fattening period in feedlot, were simulated for a 300 ha farm of the region.

Stocking rate and supplementation

The simulated system assumes that 240 kg Overo Colorado x Hereford calves are bought in April and are sold at 2 yr of age or when they attain 450 kg LW, whichever occurs first. Animals grazed during the fall and winter period, from May to September, supplemented with pasture silage. The purchase price was \$610 kg⁻¹ (1 US\$ = \$540), and the sale price is \$580 kg⁻¹ (Tattersall, 2006). Three stocking rates, 2, 2.5 and 3 head ha⁻¹, were simulated with and without grass silage supplementation at the rate of 5 kg DM d⁻¹ head⁻¹.

Each scenario was replicated 20 times. The output variables of interest included final weight and age of steers, mean total and operational cost per kg LW, the financial income and the operational income of the farm. Means were compared by the least significant difference (LSD, $P \leq 0.05$) (Table 2). In all cases, simulation runs were terminated upon reaching the maximum stipulated

Table 1. Input–output data and observed and simulated output data for five consecutive years of fattening on pasture, pasture silage conservation and supplementation. Input data and initial conditions are as reported by Rojas and Romero (1994). Simulated results are means of 20 replicates \pm SD.

Variables	Year 1	Year 2	Year 3	Year 4	Year 5	
Input	Stocking rate, steers ha ⁻¹	2.5	3.0	3.5	4.0	5.0
	Area conserved, %	26.2	14.1	12.0	3.9	0
	Initial date (day/month)	09/04	15/04	14/04	05/04	05/04
	Ending date (day/month)	10/12	14/01	05/01	23/12	18/01
	Grazing period, days	255	274	266	270	288
	Days of supplementation	-	-	31	75	92
	Initial live weight, kg steer ⁻¹	194	210	205	206	189
Observed						
Output	Final live weight, kg steer ⁻¹	378	388	390	386	342
	Pasture conserved DM, kg	8160	4668	4397	1333	-
	Simulated					
	Final live weight, kg steer ⁻¹	374.3 \pm 8.91	390.8 \pm 7.34	390.6 \pm 5.51	387.4 \pm 5.93	341.4 \pm 9.75
	Pasture conserved DM, kg	8463 \pm 943	4790 \pm 625	4392 \pm 998	1351 \pm 52	-

DM: dry matter.

Table 2. Simulated final live weights and financial performance in response to changes in stocking rate and pasture silage supplementation.

Stocking rate (steers ha ⁻¹)	Supplemen- tation (yes/no)	Final weight (kg)	Mean total cost	Mean operational cost	Total income	Operational income	Sales
		kg	——— \$Ch kg ⁻¹ ———		——— million \$Ch ———		000' kg
2.0	No	450.3 \pm 6.6c	473.6 \pm 5.5a	386.2 \pm 5.5a	23.3 \pm 1.8c	46.7 \pm 2.1bc	270.1 \pm 3.9a
	Yes	495.6 \pm 3.6f	469.9 \pm 3.0a	397.2 \pm 2.9b	26.7 \pm 1.1d	48.4 \pm 1.2c	297.3 \pm 2.2c
2.5	No	418.3 \pm 4.2b	501.3 \pm 4.5c	414.3 \pm 4.3c	18.4 \pm 2.5b	45.7 \pm 1.8b	313.7 \pm 3.2d
	Yes	493.5 \pm 3.6e	470.6 \pm 2.7a	397.6 \pm 3.1b	33.1 \pm 1.2e	60.1 \pm 1.6d	370.1 \pm 2.8e
3.0	No	307.7 \pm 13.2a	653.2 \pm 25.1d	562.7 \pm 23.7d	-25.5 \pm 5.9a	-0.5 \pm 6.6a	276.9 \pm 11.8b
	Yes	466.9 \pm 7d	492.6 \pm 6.7b	419.6 \pm 6.4c	28.7 \pm 3.2d	59.0 \pm 3.5d	420.2 \pm 6.3f

Vertical means with different letter differ significantly ($P \leq 0.01$).

age of sale of 301 d. On the other hand, significant differences were found in terms of final weights, which decreased with increasing stocking rate regardless of supplementation level, although silage supplementation tended to reduce differences between stocking rates. All associated costs increased with increasing stocking rates due to falling final weights (Figure 1). These results are in accordance with what was reported by Rojas *et al.* (2004), who used similar stocking rates at 2.5 steers ha⁻¹, but in the experiment LW gain per hectare was higher, probably due to the use of a higher amount of supplement.

Unsupplemented LW production per hectare increased initially but then fell at the highest stocking rate due to limited forage availability. When silage was provided, production per hectare increased (Table 3) although at a decreasing rate at the highest stocking rate, again due to low forage availability during the non-supplemental period.

Total and operational income initially increased with stocking rate in all simulated systems, after which the diminishing daily weight gains did not compensate for the increase in the number of animals (Table 2).

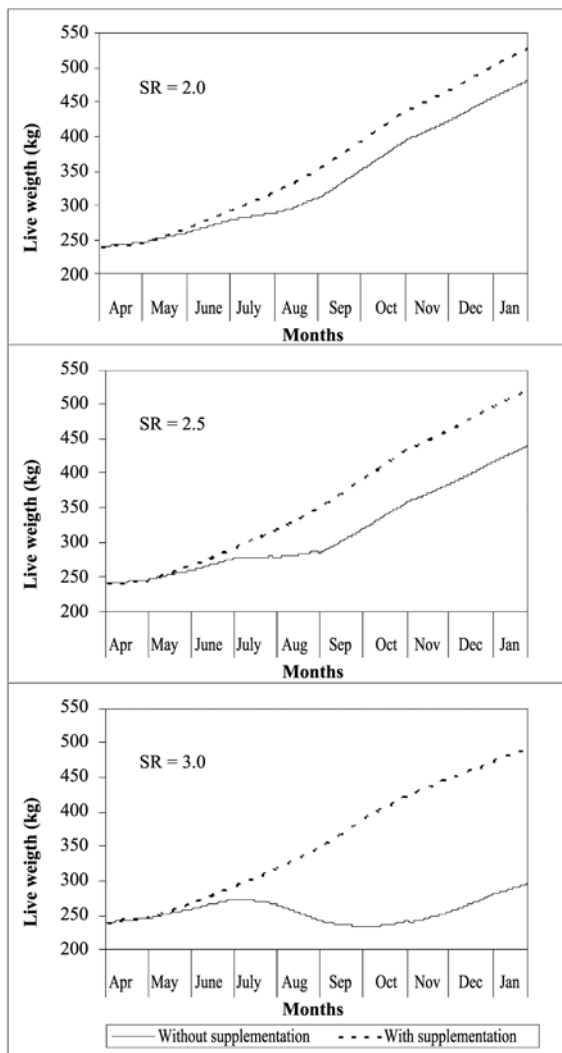


Figure 1. Simulated effects of three stocking rates (SR, steers ha⁻¹) and supplementation on the evolution of the live weight of grazing steers.

Table 3. Simulated weight gain per hectare at different stocking rates, with/without pasture silage supplementation.

Stocking rate	Not supplemented	Supplemented
steers ha ⁻¹	kg ha ⁻¹	kg ha ⁻¹
2.0	540	631
2.5	596	784
3.0	383	861

As expected, the inflexion point is reached first in the unsupplemented treatments, whereas supplementation partially compensated for the increased stocking rates.

Stocking rate, supplementation and feedlot finishing

A second set of simulation runs was carried out to simulate on-farm steer finishing through a final phase of confined feeding. This last phase lasted 80 d, and was based on a ration of silage and concentrates containing 2.3 Mcal EM kg⁻¹ DM, 14% CP and a cost of \$50 kg⁻¹ DM with the objective of achieving a final weight of 480 to 540 kg at an age about 20 months. The remaining conditions were as described before.

Age at sale was the factor determining the length of the simulated period, which in all cases lasted 380 d. As shown in Table 4, final weights differed between stocking rates and systems, but the differences are smaller than those reported in Table 1, due to compensatory gains realized during the feedlot period.

The calculated mean costs and total income (Table 4) show that the optimum is 2.5 steers ha⁻¹, since this stocking rate provides the highest income at the lowest mean cost. Comparison to the previous set of simulations shows that at stocking rates of 2 and 2.5 steers ha⁻¹, the financial income of unsupplemented systems increases by 17 and 22%, respectively, whereas supplementation allows increases of 22 and 44%, respectively.

CONCLUSIONS

The use of silage supplementation presented a positive economic and productive effect on the final live weight of the animals, regardless of the stocking rate.

Increasing the stocking rate up to 2.5 steers ha⁻¹ is feasible if winter supplementation is available, independently of a finishing period in the feedlot.

Although it is feasible to increase the stocking rate up to 3 steers ha⁻¹ with the highest live weight gain per hectare (861 kg), the weight of the animals individually will be lower because of a low availability of pasture, irrespective of the amount of supplementation.

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Table 4. Mean and standard deviation for indices of physical and economic performance of the system as a function of stocking rate and supplementation.

Stocking rate	Supplementation	Final weight	Total mean cost per kg live weight	Operational mean cost per kg live weight	Total income	Operational income	Live weight sales
Steers ha ⁻¹	yes/no	kg steer ⁻¹	————	\$Ch ————	—— million	\$Ch ————	000 ³ kg
2.0	no	496.7 ± 6.6b	521.5 ± 3.4b	433.5 ± 5.8a	29.1 ± 2.0b	55.3 ± 2.5a	298.0 ± 4.0a
	yes	540.4 ± 3.9e	517.8 ± 2.7a	440.7 ± 3.2b	32.8 ± 1.1c	57.8 ± 1.5b	324.2 ± 2.4b
2.5	no	468.8 ± 5.4a	543.2 ± 5.0c	457.4 ± 5.2d	26.6 ± 2.1a	56.8 ± 2.4b	351.6 ± 4.0c
	yes	533.4 ± 4.5d	521.6 ± 5.4b	445.0 ± 3.7c	38.9 ± 1.7e	69.6 ± 2.1c	400.0 ± 3.4d
3.0	no	i	i	i	i	i	i
	yes	505.7 ± 5.3c	543.9 ± 5.2c	468.6 ± 5.0e	34.2 ± 2.7d	68.4 ± 3.0c	455.1 ± 4.8e

Vertical means with different letters differ significantly ($P \leq 0.01$): i The system is infeasible without supplementation.

RESUMEN

Modelación de estrategias de suplementación en la recría y engorda de novillos en el Sur de Chile. Un modelo matemático fue desarrollado para analizar sistemas de producción de carne bovina en el Sur de Chile. El estudio consideró la identificación de los componentes en diferentes estrategias usadas en novillos de carne, usando praderas con y sin suplementación y la recría seguida por una engorda a corral en invierno con ensilaje de praderas y granos. La validación de los resultados a pradera del modelo usando novillos Hereford de 200 kg PV contra resultados experimentales no mostró diferencias significativas ($P \geq 0.01$) entre los pesos finales simulados y los observados. En el estudio, fueron simuladas tres carga animal (CA) de 2, 2.5 y 3 novillos ha⁻¹ con y sin suplementación de ensilaje de pradera a una tasa de 5 kg MS novillo⁻¹ día⁻¹ durante todo el período de alimentación. Las medias fueron comparadas por Diferencias Significativas Mínimas (LSD, $P \leq 0.05$). Diferencias significativas se encontraron en los pesos finales, que disminuyeron en razón del aumento de la CA independiente del nivel de suplementación, aunque ésta tendió a reducir las diferencias entre las CA. Se simuló además la respuesta de los novillos en el período final de engorda a corral y con una ración base de ensilaje y granos. Los pesos finales difirieron entre CA y los resultados mostraron que la CA de 2.5 novillos ha⁻¹, entregó el mayor ingreso al menor costo medio. Se concluye que una carga animal de 2.5 novillos ha⁻¹ es factible si existe disponibilidad de suplementación, independiente del sistema a corral para la terminación.

Palabras clave: bovinos de carne, suplementación, pastoreo, modelos de simulación.

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A STUDY OF DAIRY FARM TECHNICAL EFFICIENCY USING META-REGRESSION: AN INTERNATIONAL PERSPECTIVE

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ABSTRACT

This paper develops a meta-regression analysis to explain the variation of mean technical efficiency (PETP) measurements from a total of 65 frontier studies that report technical efficiency (ET) measurements at the dairy farm level in the literature published in English and Spanish. The analysis includes the effect of methodology on ET measurements, as well as the effect of the econometric procedure on the meta-regression estimates. Eight models were estimated, and two of these were selected: a fixed effects specification with dummy variables for the most significant studies without geographical effects (EFS), and a specification where the multiple observations are averaged and geographical effects included (OP). Based on model performance, the EFS option is chosen for the analysis. The results of the EFS model suggested that non-parametric deterministic models generate higher PETP estimates than the parametric cases (stochastic and deterministic frontier models). In addition, the Cobb-Douglas and translog forms yield higher average PETP than all other functional forms, cross-sectional data produce higher ET estimates than panel data, and the PETP is higher when the study is input-oriented. The primal approach implies a higher ET estimate than the dual analysis, and when more variables are included in the model, the PETP value is higher.

Key words: meta-regression, frontier models, technical efficiency, dairy farms.

INTRODUCTION

In an environment of growing liberalization, productivity growth, which is a major element of competitiveness, is essential to insure the prosperity of agriculture in general and dairy farming in particular (Sandrey and Scobie, 1994; Pinstrup-Andersen, 2002; Ruttan, 2002). A clear example is New Zealand, which opened its economy to the world market at the beginning of 1984 and then experienced a clear improvement in farm technical efficiency (ET henceforth) (Sandrey and Scobie, 1994; Evans *et al.*, 1996; Paul *et al.*, 2000). This improvement in ET has occurred as New Zealand has experienced a marked increase in the value of dairy products exported (Blayney and Gehlhar, 2005). The measurement of ET is important because it can help in both policy formulation and farm management (Russell and

Young, 1983; Kalirajan, 1984; Bravo-Ureta and Rieger, 1991). Producers benefit directly from improvements in their technical performance because more efficient farms tend to generate higher incomes and thus have a better chance of surviving and staying in business (Bravo-Ureta and Rieger, 1991; Dartt *et al.*, 1999; Lawson *et al.*, 2004).

In the past decades, many researchers have developed and applied diverse methods to evaluate ET at the farm level. Battese (1992), and Bravo-Ureta and Pinheiro (1993) reviewed selected articles in order to derive general conclusions about the range of ET and the performance of the methodologies reviewed. Rivas (2003) applied a meta-regression analysis to describe the behavior of ET for a limited group of dairy farm studies listed in selected databases in the English language literature. More recently, Bravo-Ureta *et al.* (2007) developed a meta-regression analysis of ET measures for all agricultural activities which includes 167 farm level studies from around the world.

In this paper we contribute to the existing literature by undertaking a meta-regression analysis focused on dairy farm ET. Thus, we examine the impact of various attributes of a dairy efficiency study (e.g., estimation technique, functional and sample size, among others)

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on ET estimates. In our analysis we also account for the possible lack of data independence stemming from the presence of multiple observations from the same study.

MATERIALS AND METHODS

Data collection

A systematic search was made for dairy farm studies published in both English and Spanish between January 1986 and January 2006 in the following databases: Agricola; Agris International; Econlit; Factiva; Infotrac; Ingenta; JSTOR; ProQuest; Social Science Citation Index; Science Direct; Web of Knowledge; Web of Science; and the World Agricultural Economics and Rural Sociology Abstracts. A complementary search was performed in the following web databases: Blackwell Synergy; EconPapers; Scielo; SpringerLink; and Taylor & Francis. In addition, a search was performed in the following Spanish language literature sources: Dialnet (online database); Agrociencia (Mexico and Uruguay); Ciencia e Investigación Agraria (Chile); Cuadernos de Economía (Chile and Colombia); Economía Agraria y Recursos Naturales (Spain); Estudios de Economía Aplicada (Spain); Investigación Agraria, Producción y Sanidad Animales (Spain); Producción Animal (Spain); Revista Brasileira de Economía (Brazil); Revista de Análisis Económico (Chile); Revista Española de Estudios Agrosociales y Pesqueros (Spain); and Revista de Estudios Agrosociales (Spain).

Variable Definition and Empirical Models

The frontier function methodology, as introduced in the path breaking paper published by Farrell just over 50 years ago (1957), uses the efficient unit isoquant to measure economic efficiency (EE), and to decompose this measurement into ET and allocative efficiency (AE). In this model, ET is defined as the ability of the firm to produce maximum output given a set of inputs and the technology. AE measures the success of the firm in choosing the optimal input proportions, i.e., where the ratio of marginal products for each pair of inputs is equal to the ratio of their market prices. In Farrell’s framework, EE is a measurement of overall performance and is equal to ET times AE ($EE = ET \times AE$). These concepts are illustrated in Figure 1, where point P represents an inefficient firm and the distance QP is the amount by which all inputs could be reduced (proportionally) without lowering output to achieve the technically efficient level of production (point Q). Thus, the ET measurement is equal to the ratio $0Q/0P$. Similarly, AE is equivalent to the ratio $0R/0Q$.

The working hypothesis of this paper is that the variation in average farm ET (PETP henceforth) for dairy farms in published studies can be explained by the major attributes of the models used. For this purpose, the

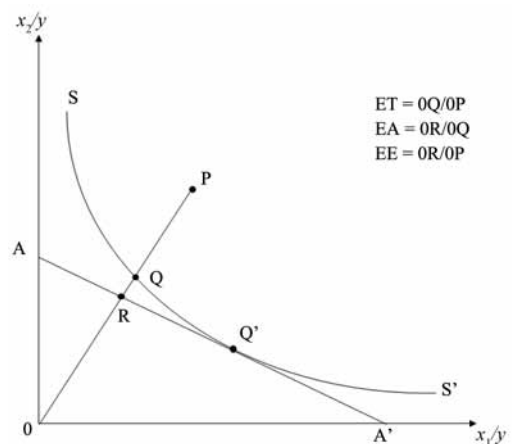
following two base models are estimated:

$$\text{Base Model A: PETP} = f(\text{PEST, PDET, TL, CD, CTR, PROD, PRI, VAR, VAROBS}) \quad [1]$$

$$\text{Base Model B: PETP} = f(\text{Model A, plus INDIA, NAMR, AFRI, LATIN, ESTE}) \quad [2]$$

The dependent variable in the meta-regressions is the PETP measurement reported in the studies included in the data set. The independent methodological variables are: PEST, a dummy equal to one if the model is a parametric stochastic frontier, and zero otherwise; PDET, a dummy equal to one if the model is a parametric deterministic frontier, and zero otherwise, the omitted category being non-parametric deterministic studies; TL, a dummy equal to one if the TL functional form is used; CD, a dummy equal to one for the CD functional form Cobb-Douglas (CD) and the excluded category is other functional forms and non-parametric studies; CTR, a dummy equal to one if the data is cross-sectional, and zero if panel data; PROD, a dummy equal to one if the model is output-oriented, and zero if input-oriented; PRI, a dummy equal to one if a primal model is estimated, and zero for dual models; VAR, the number of explanatory variables; and VAROBS, the ratio between VAR and the number of observations used in a study.

In Base Model B, the following set of regional variables is incorporated: INDIA, which is a regional dummy variable equal to one if the study used data for that part of the world, and zero otherwise; NAMR, a dummy equal to one if the data comes from North America (United States and Canada), and zero otherwise; AFRI, a dummy equal to one if the study used data from Africa, and zero



Source: Coelli *et al.* (2005).

Figure 1. Technical (ET), allocative (EA) and economic efficiency (EE) for an input oriented model.

otherwise; LATIN, a dummy equal to one if the study used data from Latin America, and zero otherwise; and, ESTE, a dummy equal to one if the study used data from Eastern Europe, and zero otherwise. The omitted regions are Western Europe and Oceania.

Meta-studies often incorporate articles that include several observations, which gives rise to a potential lack of independence in the data because studies with a higher number of observations have more weight in the analysis (Anderson and Weitz, 1989; Van Den Bergh *et al.*, 1997). Several econometric procedures have been proposed to deal with this issue. Phillips (1994) applied fixed effects, while Verlegh and Steenkamp (1999) used a two step process following a procedure suggested by Anderson and Weitz (1989). Another approach is to average the data according to some specified criteria (Espey *et al.*, 1997; Verlegh and Steenkamp, 1999; Johnston *et al.*, 2003; Hunter and Schmidt, 2004). Including a dummy variable to capture the study (fixed) effect to address the multi author problem (Anderson and Weitz, 1989) has been criticized by Verlegh and Steenkamp (1999), who argue that incorporating study dummies in a meta-regression model is likely to introduce severe multicollinearity. To avoid the collinearity problem, Anderson and Weitz (1989) suggest a two step procedure. First, the model without study dummies is estimated and the residuals from this step are used as the dependent variable in a second regression. In this second step, the study dummies are regressed on the residuals from the first step using a stepwise procedure. If the residuals are "white noise" then there are no study effects, and if not, then the selected dummies are introduced into the original model, which is re-estimated.

Another problem that arises when studies have multiple estimates, and thus the observations lack independence, is a possible bias in the standard errors of the meta-regression parameters, which would invalidate tests of hypotheses (Espey *et al.*, 1997; Verlegh and Steenkamp, 1999; Johnston *et al.*, 2003; Hunter and Schmidt, 2004). One option to mitigate this problem is to average multiple observations from a given study. This can be done in various ways (Hunter and Schmidt, 2004). In this paper, the presence of multiple PETP is due to the diversity of attributes used in the estimation of ET and all the main attributes are included in the base Models A and B. However, some attributes are incorporated in only a few models within a study and in such cases we average the respective ET measurements.

To deal with the various issues discussed above, three additional models (1, 2 and 3) are estimated for each base Model (A and B), yielding a total of eight estimated models: 1) Models with full fixed effects; 2) Models with selected fixed effects; and 3) Models with averaged

multiple observations. Models with full fixed effects and with selected fixed effects include a set of dummy variables that are defined for each study that reports two or more PETP estimates. Each study dummy is equal to 1 for all the observations that belong to a given study, and zero otherwise. Models with full fixed effects include all study dummies available, while models with selected fixed effects include only the selected study dummies following the two step procedure suggested by Anderson and Weitz (1989), as previously detailed.

ET scores are bounded between zero and one; thus, the two-limit Tobit procedure should be used (Greene, 2003). However, the meta-analysis literature focusing on ET in the agricultural sector reports similar results for the Tobit and Ordinary Least Square (OLS) procedures. Another consideration articulated by Stanley and Jarrell (1989) is that meta-regression studies use different data sets, different sample sizes, and different independent variables, which suggest that the variances of the meta-regression coefficients may not be equal, which implies that meta-regression errors are likely to be heteroskedastic. Therefore, in the current study all meta-regressions are estimated using White's heteroskedastic consistent covariance matrix estimation to correct the estimates for an unknown form of heteroskedasticity. This procedure is readily available in the Shazam Econometrics Software (Whistler *et al.*, 2001) and has been used in other meta-analysis work (Johnston *et al.*, 2003; Bravo-Ureta *et al.*, 2007).

RESULTS AND DISCUSSION

The literature search generated a total of 65 published papers which contain the type of information required for the present research. Because many of the papers report multiple ET estimates, the meta-dataset consists of a total of 329 observations. Table 1 presents an overview of all papers used in this assessment, including the authors, year of publication, country, and the PETP reported. In addition, all these papers are classified by the methodology implemented in the studies. To simplify the table, for studies that report more than one estimate using the same methodology, the average figures are included.

Table 2 presents the methodological features of the studies included in this research. As indicated, a total of 65 studies are included out of which 38 apply deterministic models and 33 stochastic models. It is important to mention that the total number of papers with stochastic and deterministic models (71) is larger than the reported number of papers (65) because in some studies both techniques are implemented. All studies combined yield a total of 329 observations given that, as already stated, some authors report multiple estimates. The data show a similar number of observations and studies that use

Table 1. Overview of empirical studies of average mean technical efficiency (PETP) for dairy farms.

Author(s). (Year). Journal, Country¹	Number of measurements²	Sample size (Number of farms)	PETP (%)
I. No-Parametric			
Deterministic frontier			
Arzubi and Berbel (2001), Rev. Esp. Estud. Agrosoc. Pesq., Argentina	3	35	77.8
Arzubi and Berbel (2002), Invest. Agrar. Prod. Sanid. Anim., Argentina	6	42	87.5
Arzubi <i>et al.</i> (2004), Rev. Argent. Econ. Agrar., Argentina	1	45	90.5
Asmild <i>et al.</i> (2003), J. Prod. Anal., Netherlands	2	1808	80.5
Cloutier and Rowley (1993), Can. J. Agric. Econ., Canada	2	187	89.8
Fraser and Cordina (1999), Agric. Syst., Australia	6	50	88.5
González <i>et al.</i> (1996), Invest. Agrar. Econ., Spain	8	56	77.9
Jaforullah and Whiteman (1999), Aust. J. Agric. Resour. Econ., New Zealand	1	264	89.0
Kaliba (2004), Q. J. Int. Agric., Tanzania	8	240	75.9
Lachaal <i>et al.</i> (2002), Mediterr. J. Econ. Agric. Environ., Tunisia	1	17	68.0
Mathijs and Vranken (2001), Post Communist Econ., Hungary	3	26	42.3
Pardo <i>et al.</i> (2002), Empir. Econ. Lett., Spain	5	38	65.2
Piesse <i>et al.</i> (1996), J. Comp. Econ., Slovenia	4	272	86.0
Reinhard <i>et al.</i> (2000), Eur. J. Oper. Res., Netherlands	8	1535	79.7
Silva <i>et al.</i> (2004), New Medit, Portugal	2	122	66.6
Tauer (1993), Agric. Resour. Econ. Rev., USA	2	395	78.3
Tauer (1998), J. Agric. Econ., USA	6	630	90.0
Thirtle <i>et al.</i> (1996), J. Prod. Anal., Slovenia	34	136	77.9
Thomas and Tauer (1994), Can. J. Agric. Econ., USA	4	125	89.2
Weersink <i>et al.</i> (1990), Can. J. Agric. Econ., Canada	1	105	94.9
Average			78.8
Stochastic frontier			
Haghir <i>et al.</i> (2004), Appl. Econ., Canada	12	1021	58.2
Average			58.2
II. Parametric			
Deterministic frontier			
Ahmad and Bravo-Ureta (1996), J. Prod. Anal., USA	5	1072	76.5
Álvarez <i>et al.</i> (1988), Rev. Estud. Agro-soc., Spain	1	154	40.0
Álvarez and González (1999), Am. J. Agric. Econ., Spain	1	410	72.0
Álvarez and Arias (2004), Agric. Econ., España	1	1176	70.0
Arias and Álvarez (1993), Invest. Agrar. Econ., Spain	1	336	73.0
Bravo-Ureta (1986), Can. J. Agric. Econ., USA.	1	222	82.2
Bravo-Ureta and Rieger (1990), J. Agric. Econ., USA	6	404	63.3
El-Osta and Morehart (2000), Rev. Agric. Econ., USA	3	679	87.0
Haghir and Simchi (2003), Empir. Econ. Lett., USA.	1	210	67.4
Hallam and Machado (1996), Eur. Rev. Agric. Econ., Portugal	3	340	66.3
Karagiannis <i>et al.</i> (2002), J. Prod. Anal., U.K.	22	2147	70.4
Lachaal <i>et al.</i> (2003), Eur. Assoc. Anim. Prod., Tunisia	1	61	75.0
Maietta and Sena (2000), Eur. Rev. Agric. Econ., Italy	1	533	55.0
Orea <i>et al.</i> (2004), J. Prod. Anal., Spain	3	445	65.9
Piesse <i>et al.</i> (1996), J. Comp. Econ., Slovenia	4	272	56.0

Continuated Table 1.

Poe and Jones (1992), <i>J. Am. Soc. Farm Manag. Rural Appraisers</i> , USA	4	675	74.8
Richards and Jeffrey (2000), <i>J. Agric. Resour. Econ.</i> , USA	1	181	94.2
Tauer and Belbase (1987), <i>Northeastern J. Agric. Resour. Econ.</i> , USA	1	432	69.3
Turk (1995), <i>Zb. Bioteh. Fak. Univ. Ljubl. Kmet. Supl.</i> , Slovenia	2	272	78.0
Average			70.1
Stochastic frontier			
Ahmad and Bravo-Ureta (1996), <i>J. Prod. Anal.</i> , USA	12	1072	81.0
Arias and Álvarez (1993), <i>Invest. Agrar. Econ.</i> , Spain	1	336	82.0
Bailey <i>et al.</i> (1989), <i>West. J. Agric. Econ.</i> , Ecuador	1	68	78.1
Battese and Coelli (1988), <i>J. Econom.</i> , Australia	2	336	70.0
Bravo-Ureta and Rieger (1990), <i>J. Agric. Econ.</i> , USA	2	404	83.9
Bravo-Ureta and Rieger (1991), <i>Am. J. Agric. Econ.</i> , USA	1	511	83.0
Brümmer and Loy (2000), <i>J. Agric. Econ.</i> , Germany	1	5093	96.0
Brümmer (2002), <i>Am. J. Agric. Econ.</i> , Germany, Netherlands and Poland	12	300	86.9
Cuesta (2000), <i>J. Prod. Anal.</i> , Spain	5	410	82.7
Dawson (1987), <i>Eur. Rev. Agric. Econ.</i> , U.K.	3	434	85.3
Dawson (1988), <i>Oxf. Agrarian Stud.</i> , U.K.	1	406	81.0
Dawson (1990), <i>Oxf. Agrarian Stud.</i> , U.K.	3	306	86.9
Dawson and Wales (1990), <i>Appl. Econ.</i> , U.K.	3	306	85.7
Dawson and Woodford (1991), <i>Oxf. Agrarian Stud.</i> , U.K.	1	918	86.0
Ghosh <i>et al.</i> (1994), <i>Forecast. Soc. Change</i> , USA	1	145	91.9
Haghir and Simchi (2003), <i>Empir. Econ. Lett.</i> , USA	1	210	83.1
Hallam and Machado (1996), <i>Eur. Rev. Agric. Econ.</i> , Portugal	1	340	88.0
Heshmati (1998), <i>Appl. Econ.</i> , Sweden	1	3979	94.5
Heshmati and Kumbhakar (1994), <i>J. Prod. Anal.</i> , Sweden	12	559	82.2
Jaforullah and Deblin (1996), <i>N. Z. Econ. Pap.</i> , New Zealand	3	264	91.9
Kumbhakar <i>et al.</i> (1989), <i>Rev. Econ. Stat.</i> , USA	6	89	72.2
Kumbhakar <i>et al.</i> (1991), <i>J. Bus. Econ. Stat.</i> , USA	9	519	73.4
Kumbhakar and Heshmati (1995), <i>Am. J. Agric. Econ.</i> , Sweden	13	4890	84.7
Lawson <i>et al.</i> (2004), <i>Livest. Prod. Sci.</i> , Denmark	2	574	94.5
Lawson <i>et al.</i> (2004), <i>J. Dairy Sci.</i> , Denmark	2	514	92.8
Mbaga <i>et al.</i> (2003), <i>Can. J. Agric. Econ.</i> , Canada	8	1143	94.8
Moreira López <i>et al.</i> (2006), <i>Arch. Med. Vet.</i> , Chile	5	92	72.2
Pierani and Rizzi (2003), <i>Agric. Econ.</i> , Italy	7	533	65.9
Reinhard <i>et al.</i> (1999), <i>Am. J. Agric. Econ.</i> , Netherlands	2	1545	89.9
Reinhard <i>et al.</i> (2000), <i>Eur. J. Oper. Res.</i> , Netherlands	8	1535	89.4
Reinhard and Thijssen (2000), <i>Eur. Rev. Agric. Econ.</i> , Netherlands	11	2589	83.8
Saha and Jain (2004), <i>Indian J. Agric. Econ.</i> , India	8	23	90.2
Average			83.3
OVERALL AVERAGE			78.4

¹ Full citations are not presented to save space and are available upon request from the authors. Journal titles are presented using ISO (International Organization for Standardization) abbreviations.

² Several studies report various measurements of ET stemming from the application of different methods.

Table 2. Summary of empirical studies of average mean technical efficiency (PETP) for dairy farms.

Category	N° Obs.	N° Studies ¹	Deterministic	Stochastic	PETP ¹
			Average (Min-Max)	Average (Min-Max)	
Approach					
Parametric	210	46	70.1 (40.0-94.2)	83.3 (47.9-99.8)	79.4
Non-Parametric	119	21	78.8 (39.0-100.0)	58.2 (42.0-69.0)	76.7
Data					
Panel	207	30	75.6 (46.0-94.2)	79.7 (42.0-99.8)	77.7
Cross Sectional	122	35	75.5 (39.0-100.0)	84.9 (47.9-96.6)	79.6
Functional form¹					
Cobb-Douglas	72	22	73.0 (40.0-94.2)	79.8 (47.9-92.5)	77.9
Translog	114	21	69.5 (49.0-85.6)	85.9 (60.9-99.8)	81.2
Others	24	5	65.6 (46.0-79.7)	81.3 (61.8-96.6)	75.4
Returns to scale					
Constant	129	39	75.3 (39.0-100.0)	76.3 (42.0-95.0)	75.8
Variable	200	38	75.7 (46.0-94.9)	85.0 (60.9-99.8)	80.1
Orientation					
Output	202	48	73.1 (40.0-94.9)	81.5 (42.0-99.8)	78.7
Input	127	26	77.2 (39.0-100.0)	80.8 (61.8-95.0)	77.9
Technology representation					
Primal	282	54	75.6 (39.0-100.0)	83.1 (42.0-99.8)	78.9
Dual	47	11	75.6 (49.0-94.2)	75.5 (47.9-88.5)	75.5
Language					
English	303	58	75.1 (39.0-100.0)	81.7 (42.0-99.8)	78.4
Spanish	26	7	79.3 (40.0-92.5)	73.8 (69.0-82.0)	78.0
Geographical region					
Africa	10	3	75.1 (58.7-86.4)		75.1
India	8	1		90.2 (86.6-92.5)	90.2
Latin America	16	5	84.9 (76.9-92.9)	73.2 (69.0-78.1)	80.5
North America ³	89	19	78.8 (45.9-100.0)	75.9 (42.0-96.6)	77.1
Eastern Europe	47	4	74.5 (39.0-93.0)		74.5
Western Europe and Oceania	159	33	73.2 (40.0-90.8)	84.2 (60.9-99.8)	79.7
Total average			75.6 (39.0-100.0)	81.4 (42.0-99.8)	78.4
Number of observations			169	160	329
Number of studies²			38	33	65

¹ Valid for parametric approach only.² Several studies report various measurements of ET stemming from the application of different methods.³ North America includes the USA and Canada.

deterministic (169 observations) and stochastic models (160 observations). The PETP for all deterministic models is 75.6% compared to 81.4% for all stochastic models and this mean difference is statistically significant at 5%. In addition, most of the studies rely on the translog (TL) functional form, are output-oriented and are mainly published in English (58 out of 65).

Table 2 also summarizes the PETP measurements according to the geographical region where the studies were

conducted. Western Europe and Oceania have the largest number of observations (159 in 33 studies), followed by North America (89 in 19 studies), Eastern Europe (47 in four studies), Latin America (16 in five studies), Africa (10 in three studies) and India (eight in one study). The highest PETP, when stochastic and deterministic studies are combined, is for India (90.2%), while the lowest is for Eastern Europe (74.5%).

Table 3. Meta-regressions of mean technical efficiency (PETP) for dairy farms.

Variables	Selected fixed effects (EFS)	Averaged model (OP)
Constant	67.524 *** 3.520 ^a	72.169 *** 4.725
PEST, parametric stochastic frontier	-6.857 4.256	2.908 5.006
PDET, parametric deterministic frontier	-18.855 *** 3.994	-9.098 * 4.922
TL, translog	13.059 *** 4.469	2.770 5.575
CD, Cobb-Douglas	15.117 *** 4.244	2.653 4.834
CTR, cross-sectional	2.426 ** 1.213	-1.706 2.471
PROD, output-oriented	-2.456 * 1.256	-3.688 2.852
PRI, primal model	9.137 *** 2.968	4.673 3.449
VAR, number of explanatory variables	0.240 *** 0.082	0.264 * 0.150
VAROBS, ratio between VAR and the number of observations	3.546 2.571	-5.107 7.681
INDIA, India		14.378 ** 6.721
NAMR, North America		6.244 ** 2.821
AFRI, Africa		0.535 5.866
LATIN, Latin America		5.595 4.277
ESTE, Eastern Europe		-9.700 ** 4.380
Log-likelihood	-1.098.5	-450.7
R²	0.6754	0.3726
Adj. R²	0.6329	0.2897

*** Significant at 1%; ** Significant at 5%; * Significant at 10%.

^a Figures in italics are robust standard errors.

PEST, dummy used if the model is a parametric stochastic frontier or not; PDET, dummy used if the model is a parametric deterministic frontier or not; TL, dummy used if the TL functional form is used; CD, dummy used for the CD functional form or not; CTR, dummy used if the data is cross-sectional or not; PROD, dummy used if the model is output-oriented or not; PRI, dummy used if a primal model is estimated or not; VAR, the number of explanatory variables; VAROBS, the ratio between VAR and the number of observations used in a study; INDIA, regional dummy variable if the study used data for that part or the world or not; NAMR, dummy used if the data comes from North America (United States and Canada) or not; AFRI, dummy used if the study used data from Africa or not; LATIN, dummy used if the study used data from Latin America or not; and ESTE, dummy used if the study used data from Eastern Europe or not.

A preliminary analysis reveals that the two preferred options are the Selected Fixed Effects Model (Model EFS) that includes methodological variables, without geographical variables, and the Averaged Observations Model (Model OP) that incorporates both methodological and geographical variables. These two models are not nested, so no further formal statistical comparisons among them are undertaken. The parameters for both of these models are included in Table 3 and a simple comparison of the number of significant parameters and adjusted R² reveals that model EFS is clearly superior to model OP. Therefore, the following analysis of the results is based on model EFS. Additional information for all models can be obtained directly from the authors.

The variables PEST and PDET capture the effect of the methodology used to estimate the frontier on PETP estimates where the excluded category for this group of dummies is the non-parametric approach. Model EFS has a negative parameter for PEST while in Model OP it is positive, but in both cases it is non significant. Theoretically, a positive value is expected for the parameter for PEST, given that deterministic models assume that all deviations from the frontier represent inefficiency (Coelli *et al.*, 2005). The estimated parameter for PDET suggests that parametric deterministic models yield lower PETPs than non-parametric models, which is valid in both models. This finding is also consistent with *a priori* expectations (Kumbhakar and Lovell, 2000). Thiam *et al.* (2001) found a negative and significant parameter for stochastic models compared to deterministic models in their research using 34 studies covering only developing countries. Bravo-Ureta *et al.* (2007) found a negative and significant parameter for both the parametric stochastic and deterministic models when compared with the non-parametric approach in their research using 167 studies on farming.

The TL and CD specifications are statistically significant in Model EFS, but not for Model OP. The CD and TL yield higher PETPs than other functional forms. These results suggest that the functional form has an unclear effect on PETP, which is consistent with what has been reported by Ahmad and Bravo-Ureta (1996), Resti (2000), and Bravo-Ureta *et al.* (2007), among others.

The parameter for CTR (Cross Sectional data) is positive and significant in Model EFS, which is consistent with the averages shown in Table 2, while the PROD parameter (orientation of the model) is negative. Thus, these findings suggest that frontier models using an output-oriented approach produce lower PETP estimates than models based on an input-oriented approach. Neither Thiam *et al.* (2001) nor Bravo-Ureta *et al.* (2007) include this variable in their meta-regressions.

Model EFS has a positive parameter for PRI, suggesting that the question of whether the model relies on a primal (PRI) or dual representation of the technology can have a significant effect on PETP. By contrast, Bravo-Ureta *et al.* (2007) found a non-significant effect for this variable.

The results indicate that the parameter for VAR (number of explanatory variables) is positive and significant and VARSIZE (ratio between the number of explanatory variables and the number of observations) is also positive but not significant. Thomas and Tauer (1994) reported an increase in the ET measurements in a non-parametric analysis when the number of variables is increased, which is consistent with the Bravo-Ureta *et al.* (2007) findings. In general and as would be expected, these results indicate a positive association between PETP and model dimensionality (Chavas *et al.*, 2005).

CONCLUSIONS

The empirical and the conceptual literature contain mixed results and contradictory views concerning the virtues of the various methodologies that have been developed to measure technical efficiency. This paper organizes studies originating from an extensive body of literature that has been published in English and Spanish over the past few decades on dairy farm ET. A total of 65 studies that use frontier models report PETP measurements at the farm level, and all the variables required for the estimated models are included. These studies yielded 329 observations, given that some report several PETP estimates.

Eight alternative models were estimated and several tests indicate that two of them perform better than the rest and thus are selected for further analysis. These two models are the selected fixed effects (model EFS) and the averaged multiple observations (model OP). Further analysis of the performance of these two models indicates that the EFS model is superior to the OP model. Thus, the results confirm the importance of considering the effect of multiple observations in the estimation of a meta-regression analysis.

The main results of the EFS model suggest that non-parametric deterministic models generate higher PETP estimates than the parametric cases (stochastic and deterministic frontier models). Within the parametric studies, the deterministic approach produces lower ET figures than the stochastic approach. The effect of functional form on ET is significant and the CD and TL forms yield higher average ET than all other functions. Frontier models based on cross-sectional data produce higher estimates than those based on panel data. In addition, the orientation of the study (input or output) has a

significant effect, with a higher PETP measurement being found for the input-oriented cases. The primal approach implies a higher ET estimate than the dual analysis. Finally, the dimensionality of the model is relevant and when more variables are included in the model, a higher PETP value is reported.

RESUMEN

Un estudio de eficiencia técnica en lecherías usando meta-regresión: Una perspectiva internacional. El objetivo de este estudio es realizar un análisis de meta-regresión para explicar la variación en el promedio de eficiencia técnica predial (PETP) en 65 estudios, en la literatura en inglés y español, desarrollados con datos a nivel predial y que reportan medidas de eficiencia técnica (ET). El estudio analiza tanto el efecto de la metodología empleada en la medición de la ET como el procedimiento econométrico en la estimación de la meta-regresión. Se estimaron ocho modelos de los cuales se escogieron dos: efectos fijos seleccionados (EFS) que incluye variables metodológicas y variables dummy para los estudios más significativos, y observaciones promediadas (OP) que contiene tanto variables metodológicas como geográficas. Basado en su comportamiento, se eligió el primer modelo para el análisis. Los resultados del modelo EFS sugieren que las fronteras determinísticas no-paramétricas generan PETP más altos que las paramétricas estocásticas y determinísticas. Las formas funcionales Cobb-Douglas y translogarítmica generan PETPs más altos que otras formas funcionales, datos de corte transversal producen valores de ET más altos que los de panel, y el PETP es más alto cuando el estudio es orientado al insumo. Análisis basados en el primal revelan valores promedios de ET más altos que en el dual, y un mayor número de variables incluidas en el modelo implica un PETP mayor.

Palabras clave: meta-regresión, modelos de frontera, eficiencia técnica, lecherías.

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ADMINISTRATION AND MANAGEMENT OF IRRIGATION WATER IN 24 USER ORGANIZATIONS IN CHILE

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ABSTRACT

Approximately 85% of the water consumed in Chile is destined to agricultural irrigation and is managed by the users themselves. This study analyzed the price that irrigation water users pay to their Water User Associations (WUAs) to which they belong and the relationship of this price to the professional level and performance of the WUAs. The study included 24 WUAs: 10 River Administration Boards (JV) and 14 Irrigation Canal Associations (AC). The annual operational budget of each WUA, the price paid by users and the management capacities of the board of directors of each WUA were compared. The study also determined the relative value of user payments to WUAs as a proportion of total production costs of the main crops in each zone. The variability of user fees per irrigated hectare decreases when the irrigation area of the WUA is more than 10 000 ha, though this was not observed in JVs. The presence of technical-professional staff directly affects the development and growth of the WUAs. As well, the WUAs with a greater level of capacity development (NDC) have more board members with a higher education level and have lower rates of unpaid user fees. The price that users pay to the WUA by irrigated hectare represents less than 4.0% of the average total production cost of the main crops in the study area. Finally, no correlation was found between the prices that users pay and the average profitability of the main crops, or between price and the geographical location of the WUAs.

Key words: water cost, water users associations, WUAs, water canal associations, river administration board.

INTRODUCTION

In different countries in Latin America and the Caribbean there are efforts to encourage changes in legislation and organizations related to the management and use of water for irrigation. These reforms vary from one country to another in terms of their execution, progress and content (Garduño, 2003, Huamanchumo *et al.*, 2008). Brazil, Chile, Colombia, Jamaica and Mexico have reformed the institutional character of water management, while the majority of other countries are in the process of making legal and institutional changes (Jouravlev, 2001).

Approximately 85% of water consumed in Chile is used in agricultural irrigation (Ministry of Public Work-General Water Directorate, 1999), a resource that is administered by the users themselves through organizations recognized

by law and with their own regulations and autonomous boards of directors.

The costs assumed by users for irrigation water services in the majority of Latin American countries have not been analyzed in-depth. In fact, some works have been carried out in recent years related to valuing irrigation water that tangentially address the cost of the service. Notably, there are works to estimate the economic and/or environmental value of irrigation water by Dinar (2000) cited by Çakmak *et al.* (2004), Herrera *et al.* (2004), Garrido *et al.* (2007) and Huamanchumo *et al.* (2008).

Water resources in Chile are the responsibility of the General Water Directorate, a body under the Ministry of Public Works, which grants water use rights without cost for the user. Nevertheless, the catchment, conduction and distribution of water resources, which imply administrative and operational costs, are the responsibility of Water User Associations (WUAs). These are regulated by the Water Code and have the objective of administering water sources and works through which water resources are extracted, captured and/or conducted and distributed among users. As well, they are responsible for resolving conflicts among such users (Galaz, 2004).

The Water Code (Republic of Chile, 2006) defines three types of organizations: (1) River Administration

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Boards (JV), (2) Irrigation Canal Associations (AC) and (3) Water Communities. JVs are constituted on the basis of natural water sources, including ground water. ACs are constituted on the basis of artificial water sources, normally canal networks of multi-farm irrigation systems. Water Communities are constituted on the basis of artificial water sources and distribute water according to the user rights of their members.

According to the CPA-SIGIRH-V8.03 Information System of the Public Water Registry of the General Water Directorate, there are currently 39 JVs and 168 ACs registered, which in general have professionalized their administration and comply with the functions they are charged with under existing legislation. At the same time, there are 3218 water communities registered at the national level, the majority of which are weak in technical, legal, administrative, financial and operational aspects, and with low levels of management and serious problems in intake, conduction and distribution of water resources. They have not developed institutional networks and are characterized by high levels of conflict (Melo *et al.*, 2005). As well, there are a number of additional organizations, especially among water communities, which operate informally and do not fully exercise their legal rights (Galaz, 2004).

There are WUAs in Chile that administer water for large areas under irrigation (> 30 000 ha) without having permanent engineers or technical staff, and at the same time there are others smaller WUAs with one or more professional staff, without a clear or direct relationship between the fees user have to pay to their organization (Melo *et al.*, 2005). The success of the latter has been based mainly on the capacity to professionalize the administration of the organization. As well, the support team (engineers, lawyers, technicians and accountants) carries out activities on behalf of the organization's board of directors, such as improving the quality of the service, earning extra incomes and implementing systems of gradual penalties to reduce the rate of unpaid user fees and other practices against the interests of the users.

Given this, it is estimated that a small WUA, without professional support would have difficulty in meeting all its obligations. This suggests that there is a minimum irrigation surface area under the responsibility of a WUA that makes it economically viable for the WUA to professionalize itself, taking into consideration agro-climatic and sociocultural factors.

The objective of this work was to analyze the cost that users should pay based on administration, operation and management in 24 WUAs and the relationship to the level of professionalization, management capacity and performance of these organizations and their impact on production costs.

MATERIALS AND METHODS

Twenty-four WUAs (JV and AC) from the Coquimbo to the Bio-Bio Regions were studied. The WUAs studied were willing to share their financial records, budget for the irrigation season of 2005-2006 and information about the total surface area under irrigation that they administer, as well as geographic maps and information about the administrative characteristics of the organization. The initial stage of the study was interviews with representatives of each organization (members of boards of directors, managers or administrators) with the objective of obtaining primary information. As well, a semi-structured interview was applied to the members of the boards of directors with the objective of relating their characteristics of age, years in the position, participation, contact with other organizations and educational level, among others, that can influence the functioning of the WUA. Finally, secondary information was evaluated to determine if there was a correlation between the profitability of the crops that predominate in the areas of jurisdiction of these organizations and the price that users pay for the administration of the irrigation system. The research stage and information gathering was carried out from April to June of 2006.

The level of prices and costs presented by the WUAs were evaluated, for the period of May 2005 to April 2006. The exchange rate used of \$541.7 Chilean pesos per US dollar was estimated based on data published by the Central Bank of Chile (2008), considering the average rate of exchange for the period indicated, which presented a standard deviation of \$24.2 pesos.

Primary information

This refers to the economic resources used during the irrigation season of 2005-2006 in each of the organizations evaluated, identifying or recovering administrative and operational costs, highlighting among others the operation of irrigation intakes structures, cleaning canals and professional or administrative fees. As well, this includes information related to technical and administrative aspects of the organization such as the total area under irrigation, the number of water right allotments under the jurisdiction and water resources that are managed.

The information related to the leadership capacity and performance of the board of directors of each organization was gathered by means of semi-structured interviews conducted with members of 19 of the 24 boards of the WUAs that agreed to participate in this part of the study.

Secondary information

This refers to bibliographic information available on the surface area destined to the main crops in the areas of the study (Larrañaga *et al.*, 2003; 2004; 2005; 2006), their

proportional participation in each WUA and information about the total production costs and average gross margin (Fundación Chile, 2007).

Characterization of the WUAs

To characterize the WUA an instrument termed the Capacities Development Level (NDC) was used. It was designed by the National Irrigation Commission in collaboration with the Universidad de Concepción (de Miguel, 2005). It allows for classifying, characterizing and typifying the WUAs into seven categories according to the development of their management and administrative capacities, legal situation, participation of users and other organizational aspects (Table 1). Field trips were made to each WUA and members of the boards of directors, technical staff and users were interviewed and self-evaluation test (San Martín, 2007) was applied.

Calculating the annual budget

Subsequent to the information gathering stage, a study was prepared on the operational expenses of WUAs, the resources used and administrative costs, including among others, security personnel, accountant and clerical staff, as well as the fees of professionals who render managerial-administrative and legal-technical services, infrastructural and operational expenses to maintain the functioning of the system such as upkeep of inlets and cleaning. The gathering and analysis of these data allowed for estimating the annual budget necessary for the functioning of the WUAs according to:

$$\text{Annual budget} = \text{Administration costs} + \text{Operational costs} + \text{Investments}$$

Administrative expenses include salaries (administrator, engineer, accountant, secretary, guard, benefits, bonuses, insurance and others), fees (engineer, lawyer, accountant and that are not core staff), office rental, office materials and basic services (telephone, fax, internet, electricity, water, radio), vehicle or transportation (fuel and maintenance). Operational expenses refer to the opening and closing of inlets structures, support to the JV if it corresponds, cleaning of canals, maintenance, security staffs, machine rental, trucking and transport. Investments include construction, repairs and repositioning of works and contingencies.

The total amount collected (MTR) annually by the WUA is:

$$\text{MTR} = \text{MRC} + \text{Grants} + \text{Others}$$

where MRC is the amount collected through user fees of the organization in a season (\$ season⁻¹), Grants refers to the amount obtained (\$ season⁻¹) through projects

presented under Law N° 18450 for the promotion of private investment in irrigation and drainage works, and Other considers judicial fees and extraordinary improvement programs.

Determination of cost per hectare

With the amount collected by the users fees charged to users of the system (MRC) and the total area under irrigation administered by the WUA (SBR), the price paid by user to irrigate one hectare of land (*PPU*), is expressed as:

$$PPU = \frac{MRC}{SBR} \text{ (\$ ha}^{-1} \text{ season}^{-1}) \quad [1]$$

The choice of *PPU* instead of the price for nominal flow (water right allocated as a part of the canal flow) that the user pay is due to the lack of correspondence observed between the values of the water right allotments and the water discharges that they represent, which is characteristic of and unique to each WUA (Table 2). As well, reliable water measurements are not always available throughout the irrigation seasons for all the participating WUAs, which would allow for obtaining an average seasonal water flow or fair estimation of the volume of water that corresponds to a water right allotment. Nevertheless, all the organizations have reliable and detailed information about the irrigation area under their jurisdiction, because of which, for comparative purposes, the *PPU* is adequate.

Production costs, average gross margin and price paid by users

To obtain the proportion of the cost of the administration cost of the water in the total cost of the main crops in territories of the respective WUAs, databases were reviewed with information on the main crops, based on the fruit registry of the ODEPA-CIREN (Larrañaga *et al.*, 2003; 2004; 2005; 2006). Likewise, the weighted profitability of the areas under irrigation was estimated using information on production costs, gross incomes, profitability or average gross margin (Fundación Chile, 2007), and it was determined whether there was a correlation between this parameter and the value paid by users for the administration of water resources in their system.

RESULTS AND DISCUSSION

Characterization of the WUAs

Of the 24 WUAs considered, 10 were JVs and 14 ACs (Table 2). Thus, of the total of 39 JVs and 168 ACs registered at the national level, 25.6% of the JVs and 8.3% of the ACs were included in the study. The total study represented 11.6% of the total of JVs and ACs registered in the country. Each WUA manages an irrigation system that covers more than 2500 ha, the largest being the Maule

Table 1. Level of Capacity Development (NDC) of the Water User Associations (WUAs).

Characteristics	
LEVEL OF CAPACITY DEVELOPMENT OF THE WATER USER ASSOCIATIONS	Not operative
	WUAs that do not carry out any kind of activities of their own. There can be users from some derivative administered by an Irrigation Canal Association that has the intention of forming a water community, but has not developed the necessary capacities to do so. In some case, some process of organization has been initiated (making a public document of the minutes of the first assembly), to apply for some type of grant under Law N° 18450. There are organized WUAs with their respective registries in the General Water Directorate that do not carry out any activity as a water community. In some cases, no information of the organization's legal situation, with the possibility that it forms part of another organization.
	Basic
	WUAs that carry out only basic functions, that is to say, distribute the waters of the canal network and concern themselves with canal maintenance (cleaning). They do not concern themselves with organizing the allotment of water rights, leaving the administration of this in the hands of the users, which can generate important conflicts. They do not have budgets, rarely concern themselves with improving the irrigation system that they administer, are in frequent conflicts with users, in particular in the derivatives and no effective participation is observed. In some cases users even reject the organization and do not perceive any benefits from belonging to it.
	Operative
	WUAs that carry out basic functions and concern themselves with improving existing infrastructure, for which they have developed a certain capacity of budgeting and have an annual budget that allows them to operate the systems of water intake structures, conduction and distribution. Nevertheless, there can be problems with associates not paying dues. In general the users perceive benefits from being organized. These WUAs do not organize internal information related to secondary canals, registry of users and the corresponding allotments, nor have they developed clear standards that regulate allotting water access.
	Ordered
	WUAs that are familiar with their irrigation system, can identify the secondary canals and the number of water right allotments that correspond to them. They have an organized registry of users, which facilitates charging fees. They have clearly established standards and some mechanism to resolve conflicts. They lack effective participation; many of their users only attend annual assemblies and pay their dues. No substantial renovation of the Board of Directors is observed and they lack mechanisms to improve management.
Functional	
WUAs characterized by fully meeting legal standards and by a good operation of the systems of water intake structures, conduction and distribution of available water. The users are relatively well informed about their rights and obligations and "receive the water that corresponds to them". Problems relate to the vulnerability of the system of water intake structures, conduction and distribution of water, lack of accumulation works and productive alternatives of the users.	
Dynamic	
WUAs characterized by the active participation of the users of the organization and by their capacity to take initiatives to continue strengthening it. They are capable of generating proposals and projects that allow for continued improvements in irrigation infrastructure, the internal organization and the productive possibilities of their members.	
Integrated	
WUAs characterized by having developed, as well as all the proceeding elements, effective ties with pertinent state and private bodies, ensuring an optimum use of the waters at their disposition for all their associates through a competitive and productive development based on agriculture under irrigation.	

Source: Final Report "Development of a Methodology of Organization and Training for Water Communities". Agreement between the National Irrigation Commission and the Universidad de Concepción, Faculty of Agricultural Engineering, January 2003 (de Miguel, 2005).

Table 2. Nominal discharge managed (Q_n), nominal discharge per water right allotment (Q_{Acc}) and area under irrigation (SBR) in 25 Water User Associations (WUAs) surveyed.

Region	Name of the WUA	Q_n ($m^3 s^{-1}$)	Q_{Acc} ($L s^{-1}acc^{-1}$)	SBR (Mha)
COQUIMBO	AC Punitaqui Canal	0.82	1.0	2.7
	JV Illapel River	0.40	1.0	3.5
	JV Combarbalá River	3.61	1.0	2.5
	JV Choapa River	18.26	1.0	10.7
	JV Limarí and Grande Rivers and streams	24.59	1.0	20.5
	JV Elqui River	25.34	1.0	23.5
RM	AC del Maipo	19.96	15.0	30.0
LBO	JV Cachapoal River 1 ^a Section	-	-	48.0
	AC San Pedro Población y Derivados Canal	7.36	2.7	2.7
	AC Cachapoal Canal	3.68	14.1	5.0
MAULE	JV Achibueno River	19.96	1.5	12.0
	AC Putagán Canal	4.29	1.0	20.0
	AC Maule Norte Canal	51.78	15.0	70.0
	AC Maule Sur	34.40	1.4	25.0
	JV Ancoa River and streams	7.42	1.0	9.0
	JV Maule River	199.98	1.4	180.6
BÍO-BÍO	AC del Laja	68.49	15.0	55.0
	AC Zañartu Canal	29.18	10.0	40.0
	JV Diguillín River and streams	33.39	15.1	33.0
	AC Bío-Bío Norte Canal	10.70	15.0	10.0
	AC Bío-Bío Negrete Canal	17.98	15.9	8.0
	Coihueco Reservoir Irrigators Association	4.96	1.1	5.0
	AC Duqueco Cuel Canal	7.97	1.4	6.0
AC Quillón Canal	2.29	1.0	2.5	

JV: River Administration Board; AC: Irrigation Canal Association; RM: Metropolitan Region; LBO: Libertador General Bernardo O'Higgins Region; acc: Water right allocated proportional to the river or canal discharge; Mha: Thousands of hectares.

River JV which distributes the resource to five ACs in the Maule Region, covering an area under irrigation of 180 636 ha, while the largest AC (the North Maule Canal) covers 70 000 ha under irrigation.

The 24 WUAs analyzed present various legal situations (owners of the waterway or in the process of receiving control of the canal from the State, without legal status, etc.) and diversity in terms of their organizational management. As well, there is great variability in terms of the number of water right allotments and flow discharge, surface area under irrigation that they manage, geographic location and presence of technical personnel in charge of administration. This becomes into a broad range of levels of development of capacities (NDC).

All these organizations have an administrative system and defined operational structure that coincides with its statutes, established standards and the rights and duties of the users, with a clear identification and register of its members that is periodically updated and includes, among

other things, the name, identification number, address and number of allotment holders. About 75% of the WUAs file copies of the registration of Water Use Rights in their respective Land Registry Office.

Among the WUAs analyzed there are different systems of water use rights: consumptive, permanent, and eventual rights. All of the WUAs prepare an annual budget, in which the board of directors and users participate actively in the decision making. In 100% of the WUAs the board of directors is elected, with an annual change of an average of 48% of the members. Likewise, there are annual work plans in which decisions are taken about improving infrastructure and which encourage the participation and attendance at meetings of the users. As well, in the case that the organizations have professional staff in charge of administration, training programs are carried out for members of the board of directors and for the technical and administrative teams. Finally, the NDC of the WUAs is in the range of Basic to Integrated, and

those organizations with mayor NDC in most cases have an internal accounting system, conduct administration in their own offices, have means of transport and have lower percentages of unpaid user fees, with values that fluctuate between 0 and 20% (Figure 1). These values are comparable to the 20% reported by Molle *et al.* (2008) in the Jordan Valley irrigation system, the 15% in WUAs from the coastal region of Peru (Huamanchumo *et al.*, 2008) and the 14% in WUAs in Turkey (Özlu *et al.*, 2000 cited by Çakmak *et al.*, 2004). On the other hand, the WUAs with lower NDCs have rates of unpaid user fees of up to 55%.

Price paid by the user (PPU)

In general, the ACs charge users higher prices than JVs, which is explained by the fact that JVs perform more restricted functions than ACs, limiting themselves to delivering a supply of water to the irrigation intakes of the canals under their jurisdiction. Figure 2 shows the broad variability of the PPU, oscillating between \$2665 (US\$4.9) and \$18 706 (US\$34.5) per hectare irrigated in the season, when the AC administers less than 10 000 ha. For larger surface areas, a tendency in the ACs is observed to values between \$1648 (US\$3.0) and \$7286 (US\$13.5) per hectare irrigated in the season. For the JVs, the PPU oscillates mainly between \$240 (US\$0.4) and \$3950 (US\$12.8) per hectare-season (the value of \$240 is not shown in Figure 2, and correspond to a WUA that administers 180 600 ha, which is outside the range of the graph). The exception is a JV from the Coquimbo Region, with a PPU of \$9287 (US\$17.1) per hectare, which administers a complex system of 23 500 ha and includes two water reservoirs with a total capacity on the order of 240 million m³. As well, this organization is developing a series of projects related to integrated water management and catchment, as well as education in rural schools of the province.

The PPU values obtained in this study are comparable to the price range for irrigation water per hectare, per year or season in Brazil (US\$3.5), Australia (US\$0.75 to 2.3), New Zealand (US\$6.8 to 16.6) and less than those from Mexico (US\$33 to 60), Italy (US\$21 to 78), Spain (US\$96 to 164.5), Greece (US\$92 to 210) (Dinar, 2000 cited by Çakmak *et al.*, 2004).

Professionalization of WUAs

Some 62.5% of the WUAs surveyed (15) have permanent administrative staff, while 37.5% (9) do not. Nevertheless, 91.7% (22) uses the services of an accountant, whether as permanent or on contract and 54.2% (13) have contracted and regularly draw upon professional legal services. In relation to staff for technical support (agronomists and engineers), 62.5% of the WUAs (15)

have permanent staff, 20.8% (5) contract technical support staff in the irrigation season or for specific consultations, and the rest 16.7% (4) do not have this type of support. As well, 20.8% (5) has managerial-administrative and technical-legal support that has allowed them to form parallel companies (consulting, administration or construction), responsible for managing and carrying out investment projects in the irrigation system.

Figure 3 offers a way to visualize the relationship between the degree of professionalization of the WUAs and the extra incomes obtained, where the benefits of having professional staff time are appreciable. It follows that the WUAs that contract more than 25 professional hours per week for administrative management or that are directed by an administrator who also exercises technical functions, have obtained resources that fluctuate between MM\$234 to MM\$2823. For this analysis, 22 WUAs were considered because two ACs are in the process of transferring the intake, conduction and distribution structures from the State to the users, which limits them from applying to public funding instruments.

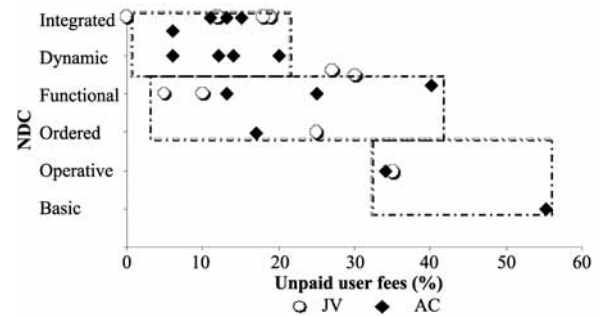
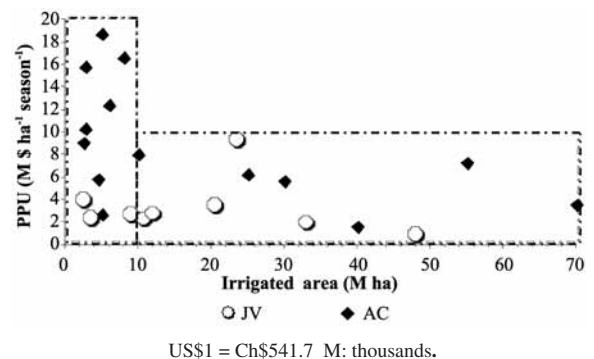


Figure 1. Relationship between the level of capacity development (NDC) and unpaid user fees in 10 River Administration Boards (JV) and 14 Irrigation Canal Associations (AC).



US\$1 = Ch\$541.7 M: thousands.

Figure 2. Price paid by users (PPU) depending on irrigated area administrated by nine River Administration Boards (JV) and 14 Irrigation Canal Associations (AC).

The percentage that the contracting of professional-administrative support of the total amount collected (MTR) by the WUAs is relatively low (Table 3). In the JVs that manage areas under irrigation greater than 30 000 ha, the cost of professional services does not exceed 16%, representing values of \$80 to \$250 ha⁻¹ season⁻¹. Nevertheless, in the JVs that irrigate from 10 000 to 30 000 ha, such costs constitute only 8%, equivalent to \$165 to \$309 ha⁻¹ season⁻¹ (San Martín, 2007).

The ACs that manage areas under irrigation larger than 30 000 ha, salaries of core staff represent 7% of the MTR in a season (Table 4), representing \$400 to \$480 ha⁻¹ season⁻¹, and in the ACs that irrigate from 10 000 to 30 000 ha and from 2 500 to 10 000 ha it represents 6% and 15%, respectively, which implies values in the range from \$400 to \$2025 ha⁻¹ season⁻¹ (San Martín, 2007).

Analysis of the interviews with members of Boards of Directors

In 19 of the 24 WUAs to which the interviews were applied 50% of the members of the board of directors are between 40 and 60 years of age and 43% are over 60. About 96% of the members of the board of directors are

male. The term of office of the board members, whether in positions as president, secretary or director, is not related to the NDC. While many WUAs establish regulations that stipulate the annual renewal of the board of directors or part of it, the fact that this change is not complied with does not mean that a WUA lacks an effective system of administration. Additionally, 52% of the WUAs that responded to the interviews do not have an annual change of the board of directors or part of it. Some larger and successful organizations prefer to conserve their directors because of the good management work they carry out. Nevertheless, other WUAs in the same category conserve part of the board of directors and carry out a partial change in accordance with the established by-law.

The directors of all WUAs with areas under irrigation larger than 10 000 ha, have some type of participation in other organizations, whether of a social or productive nature, exercising some position or participating as a member. Some of these organizations are other WUAs, fruit companies, producers' organizations or others. The smaller organizations also relate to other organizations, notably among them other WUAs, agricultural associations and neighborhood associations. It can be

Table 3. Percentage use distribution by items of the total amount collected (MTR) in 10 River Administration Boards, by range of surface area.

Item	Average of the MTR by range of surface area (ha)		
	2500-10 000	10 000-30 000	30 000 and more
Professional-administrator fees	-	8	16
Core staff salaries	39	33	15
General expenses	24	24	7
Others (investment, operational maintenance and transport, machinery rental and consulting, travel, etc.)	37	34	62
Total	100	100	100

Table 4. Percentage use distribution by items of the total amount collected (MTR) in 14 Irrigation Canal Associations, by range of surface area.

Item	Average of the MTR by range of surface area (ha)		
	2500-10 000	10 000-30 000	30 000 and more
Professional-administrator fees	15	6	7
Contracts and wages (cleaning)	29	25	39
Core staff salaries	19	32	37
General expenses	8	6	5
Others (investment, operational maintenance and transport, machinery rental and consulting, travel, etc.)	29	31	13
Total	100	100	100

deduced that in the larger organizations, the directors or users have a higher level of participation in other types of organizations, whether productive or social. Nevertheless, this indicator does not necessarily represent a better quality of management of the WUA by the directors.

The results of this study indicate that higher levels of formal education of the members of the board of directors can favor the development of the organization (Figure 4), because of which this criterion can be used as a component of the NDC of the WUAs. On the other hand, no relationship was found between the NDC and SBR administered by the WUA, nor with the renewal of the board of directors, nor with the number of water right allotments that the members of the board have.

The WUAs that have contracted professional hours show a minimum value of “Ordered” in their NDC, but when there are no professional hours contracted, the WUAs show a maximum value of “Operative” in their NDC (Figure 5). This has validity discarding an AC with zero professional hours and a “Functional” NDC, given that it received professional support for three years as part of a program for strengthening WUAs (Ministry of Agriculture-National Irrigation Commission, 2006).

Relationship between the PPU and gross margin of production

The factors that have more influence on direct production costs are inputs like fertilizers, pesticides, labor, harvesting and machinery use (Fundación Chile, 2007). By relating the value that users pay to the WUAs to the average total production cost of the main crops of the zones under study, it can be concluded that this value (PPU) represents less than 4% of the total production cost of these crops (Table 5). This value is significantly less if it is compared to the 12% average that represents the irrigation costs in two areas in India, which are derived from the information shown by Kumar *et al.* (2008)

There was no evidence of any kind of a correlation between the price paid by users (PPU) and the average gross margin (Figure 6), independent of the type of WUA or its geographic location.

CONCLUSIONS

During the 2005-2006 irrigation season, the price paid by users oscillated between \$2700 (US\$5.0) and \$18 700 (US\$34.5) ha⁻¹ season⁻¹, with considerable variability for areas of less than 10 000 ha, and values between \$1700 (US\$3.1) and \$7300 (US\$13.5) ha⁻¹ season⁻¹ for larger areas. In the JVs, the price paid by users varied between \$240 (US\$0.4) and \$9300 (US\$17.2) ha⁻¹ season⁻¹, without relating them to the area under irrigation that the WUA administers.

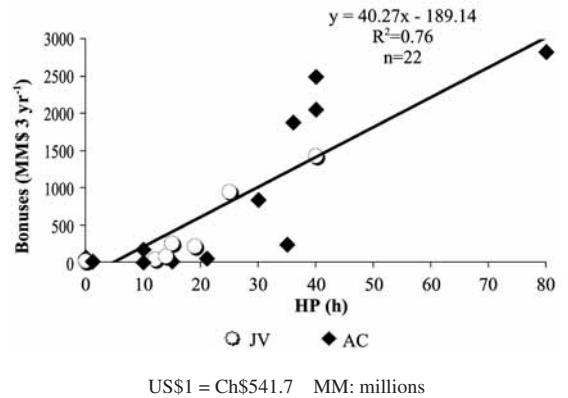


Figure 3. Extra income obtained as bonuses between 2003 and 2005 and weekly professional time (HP) hired in 10 River Administration Boards (JV) and 14 Irrigation Canal Associations (AC) (four JVs and one ACs are in the origin).

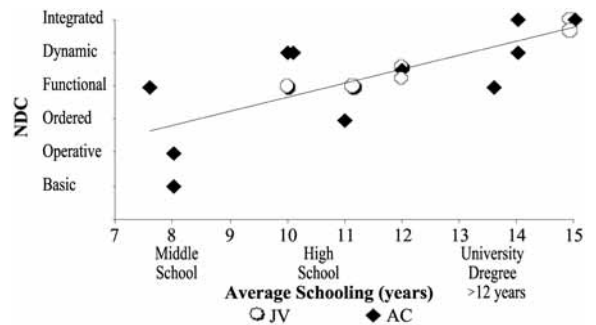


Figure 4. Relationship between Level of Capacity Development (NDC) and average schooling of the council members in seven River Administration Boards (JV) and 12 Irrigation Canal Associations (AC) surveyed.

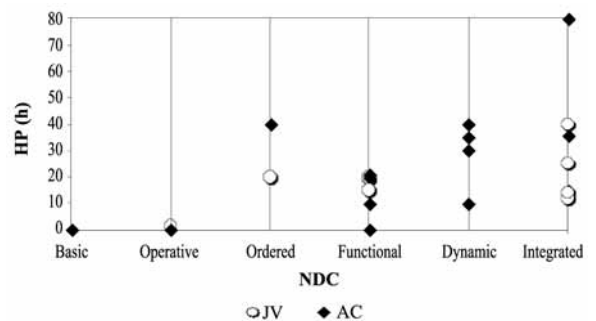


Figure 5. Weekly professional time and Level of Capacity Development (NDC) in 10 River Administration Boards (JV) and 14 Irrigation Canal Associations (AC).

Table 5. Percentage representation of the value paid by the users (PPU) of River Administration Boards (JV) and Irrigation Canal Association (AC) in the total production cost for the main crops in the irrigated areas under their jurisdiction.

Region	Water user organization	Representation PPU (%)														
		Wheat	Potato	Beets	Rice	Corn	Bean	Table grapes	Wine grapes	Red apple	Lemon	Peaches	Avocado	Raspberry	Walnut	Mandarin
COQUIMBO	AC Punitaqui Canal							1.0				1.5		1.0		
	JV Illapel River										0.2	0.2	0.2	0.2		
	JV Combarbalá River							0.2			0.3	0.4	0.4	0.4		
	JV Choapa River										0.2	0.2	0.2	0.2		
	JV Limarí and Grande Rivers and streams							0.2			0.3	0.3	0.3	0.3		0.3
	JV Elqui River							0.5			1.0	1.0	1.0	1.0		0.9
RM	AC del Maipo	1.5	0.6			0.9	0.3				0.5	0.5	0.6	0.5		
LBO	JV Cachapoal River 1ª Section	1.0				0.6	0.7	0.2	0.5	0.2						
	AC San Pedro Población y Derivados Canal	3.5				2.6	3.3	1.0	2.4	0.8						
	AC Cachapoal Canal	4.0				3.1	3.8	1.5	2.8	1.0						
MAULE	JV Achibueno River	0.6	0.3	0.1	0.7	0.5	0.6			0.2				0.1		
	AC Putagán Canal	1.3	0.7	0.3	1.5	1.0	1.2			0.3				0.3		
	AC Maule Canal	0.8	0.4	0.2	0.9	0.6	0.8			0.2				0.2		
	AC Maule Sur	1.4	0.7	0.3	1.5	1.0	1.3			0.3				0.3		
	JV Ancoa River and streams	0.6	0.3	0.1	0.6	0.4	0.6			0.1				0.1		
	JV Maule River	0.1	0.0	0.0	0.1	0.0	0.1			0.0				0.0		
BÍO-BÍO	AC del Laja	2.0	0.8	0.3		1.2	1.5									
	AC Zañartu Canal	0.5	0.2	0.1			0.4									
	JV Diguillín River and streams	0.5	0.2	0.1			0.5									
	AC Bío-Bío Norte Canal	1.8	0.9	0.4			1.7									
	AC Bío-Bío Negrete Canal	3.8	1.9	0.8		2.7	3.5									
	Asociación de Regantes del Embalse Coihueco Reservoir Irrigators Association	0.6	0.3	0.1			0.6									
	AC Duqueco Cuel Canal	2.8	1.4	0.6		2.0	2.6									
AC Quillón Canal	2.0	1.0	0.4			1.9										

Source: Own elaboration. RM: Metropolitan Region. LBO: Libertador General Bernardo O'Higgins Region.

The evaluation of WUAs involved in the study indicates that the administrative management systems differ among organizations of the same type and among the regions under study. Management capacity, measured in terms of the NDC, can be correlated with the average level of schooling of the directors of the WUAs, influencing the percentage of unpaid user fees, in the degree of professionalism or professional hours contracted and the extra incomes that are earned in the WUA. It is observed that the hiring of technical-professional staff in charge of administrative, operational and managerial tasks directly influences the development and growth of

the WUA, and provides the tools and vision necessary to obtain extra incomes. The hiring of a professional-administrator represents between 6% and 16% of the total amount collected by the organization, with lower values in WUAs that manage an area under irrigation of 10 000 to 30 000 ha.

The price paid to WUAs by users represent less than 4% of the total average cost of production of the main crops in the respective territories and are not related to the incomes generated by the producers or the geographic location of the WUA.

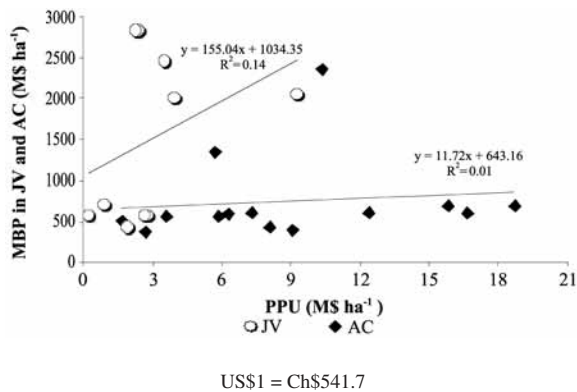


Figure 6. Average gross income margin (MBP) of the area under irrigation in 10 River Administration Boards (JV) and 14 Irrigation Canal Associations and price paid by the users (PPU) for each irrigated hectare.

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RESUMEN

Administración y gestión del agua de riego en 24 organizaciones de usuarios en Chile. Aproximadamente, el 85% del agua consumida en Chile es destinada al riego agrícola, siendo administrada por los propios usuarios. En este estudio, se analizó el costo que cancelan los usuarios del agua de riego a sus Organizaciones de Usuarios de Agua (WUAs) y el nivel de profesionalización y desempeño de éstas. Se estudiaron 24 WUAs: 10 Juntas de Vigilancia (JV) y 14 Asociaciones de Canalistas (AC). Se comparó el presupuesto anual de operaciones de cada WUA, el valor que cancelan los usuarios y las capacidades de gestión de la directiva con el desempeño de la respectiva WUA. Además, se analizó la significancia del pago de los usuarios en los costos de producción de los principales cultivos en cada zona. En las AC, la variabilidad de precios por hectárea regada que cancelan los usuarios, disminuye cuando el territorio de la WUA es mayor a 10 000 ha, situación no detectada en las JV. El personal técnico-profesional de apoyo incide directamente sobre el desarrollo y crecimiento de las WUAs. Asimismo, las WUAs con nivel de desarrollo de capacidades (NDC) más elevado poseen directivas con mayor escolaridad

promedio y los usuarios presentan menor morosidad en el pago. El precio que cancelan los usuarios a las WUAs por hectárea regada representa menos del 4,0% de los costos promedio totales de producción de los principales cultivos de cada zona. Finalmente, no se evidenció correlación entre el precio que cancelan los usuarios y la rentabilidad promedio de los cultivos, ni por ubicación geográfica de las WUAs.

Palabras clave: costo del agua, organizaciones de usuarios del agua, asociaciones de canalistas, juntas de vigilancia.

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FECAL CONTAMINATION OF GROUNDWATER IN A SMALL RURAL DRYLAND WATERSHED IN CENTRAL CHILE

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ABSTRACT

Research on microbiological groundwater quality was conducted in Chile in a rural watershed that has almost no other water source. Forty-two wells were randomly selected and levels of indicator bacteria - total coliforms (TC), fecal coliforms (FC), and fecal streptococci (FS) - were repeatedly measured during the four seasons of 2005. The aim of this study was to characterize microbiological groundwater quality, relate indicator levels to certain watershed features and management characteristics which are likely to affect water quality. The dynamics of seasonal temporal contamination was determined with statistical analyses of indicator organism concentrations. Nonparametric tests were used to analyze relationships between bacterial indicators in well water and other variables. TC, FC, and FS were found in all samples indicating the wells had been contaminated with human and animal fecal material. The frequency distribution of microorganisms fitted a logistic distribution. The concentrations appeared to be temporal and levels varied between seasons with higher concentrations in winter. The cause of contamination could be linked to the easy access of domestic animals to the wells and to the permeable well casing material. Local precipitation runoff directly influenced the bacterial concentrations found in the wells.

Key words: biological contamination, bacteria, water quality, environmental pollution.

INTRODUCTION

Water quality is a key environmental issue involving natural watershed resources and local rural communities. The major environmental pressures have an impact on the quantity and quality of groundwater resources (Danielopol *et al.*, 2003) which are generally perceived as being less vulnerable to contamination than surface water given the natural filtering ability of the subsurface. Although most groundwater is still thought to be free of disease-causing microorganisms, many systems are unprotected and contamination events could eventually occur because private groundwater wells are rarely, if ever, monitored.

The risk of contaminated water for people was manifested in Lake Erie, Ohio, USA in 2004 when 1450 people became ill because of a pathogen in the well water (Fong *et al.*, 2007). Furthermore, an estimated 750 000 to 5.9 million people are sick every year as a result of contaminated groundwater in the USA (Macler and Merkle, 2000).

One of the most frequent types of contamination in rural areas is fecal pollution from different sources, most frequently livestock and inadequate on-site human waste disposal systems (Conboy and Goss, 2001; Barnes and Gordon, 2004). The size and shape of pathogenic microorganisms, their surface density properties, and biological activities set them apart from other contaminants that are transported in surface and subsurface water environments (Pachepsky *et al.*, 2006). Concentrations of microbiological contamination indicator organisms observed in groundwater are a function of the contamination sources active at that moment (Solo-Gabriele *et al.*, 2000).

Microbiological contamination is dispersed, sporadic, and influenced by a range of interacting environmental factors such as the watershed's physical characteristics, climatic conditions, and agricultural management practices. Since the largest numbers of fecal coliforms and fecal streptococci are always present in manure

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(Chadwick and Chen, 2002), then the presence of either of these microbes in a well water sample is strong evidence of fecal contamination. One of the difficulties in tackling this problem is the fact that contamination is likely to come from various possible point and nonpoint sources (Mahler *et al.*, 2000), thus obscuring its origins. It is important to detect fecal contamination in groundwater, especially if there are no pre-consumption water treatment systems (Atherholt *et al.*, 2003). This is the case in some rural dryland areas of Chile where farmers obtain small amounts of water from private wells and face serious water supply problems for both human consumption and agricultural activities.

Improving the quality of groundwater resources offers an important economic opportunity for the gradual improvement of the quality of life in rural dryland communities. In order to develop strategies to diminish or eliminate microbiological contamination in groundwater wells, it is first necessary to assess the variability in its concentrations, and the relative importance of different factors affecting pollution.

The variability of microorganism concentrations in Chilean groundwater and the factors affecting them are not well-known at present. As rural communities continue to rely on shallow groundwater, it is important to improve the state of knowledge about the quality of this resource. To assess the presence of fecal contamination in a rural watershed, a study was undertaken to typify the quality of microbiological groundwater, describe its seasonal pattern, and look for probable characteristics exerting an influence on the quality of groundwater.

MATERIALS AND METHODS

The small rural Estero San José (ESJ) watershed (10.8 km²) is located in the Bío-Bío Region, Chile (Figure 1). The catchment area is sparsely inhabited by families dedicated to traditional agriculture. The ESJ watershed is characterized by a Mediterranean climate with a long dry season leading to water shortages and a short wet season.

The watershed soils have low permeability and capacity to provide underground water. Moisture accumulation in the watershed takes place between April and June. The major runoff period of the year is from July to October when the ground is saturated and almost all the precipitation that falls in the watershed runs off. Precipitation is scarce between November and March, with practically no base flow in the watershed. Farmers obtain small amounts of water from private wells. On the average, these are 7.0 m deep and yield a median of 1.1 L min⁻¹. Groundwater is used as drinking water, for other domestic purposes, orchards, gardens, greenhouses, and livestock production. Agricultural production in the area

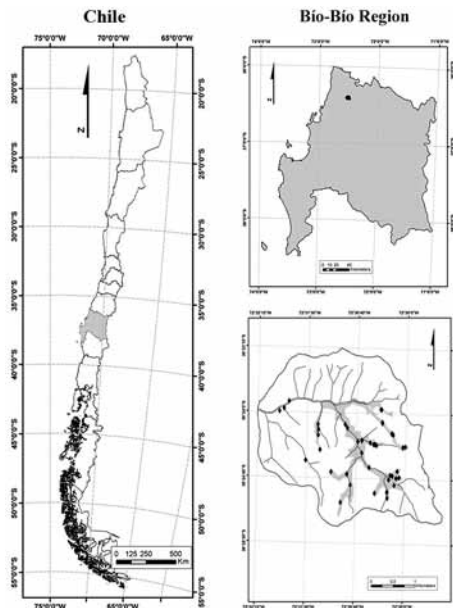


Figure 1. Location of the Estero San José Watershed and sampling sites.

is mostly wheat (*Triticum aestivum* L.) and lentils (*Lens culinaris* Medik.). The density of domestic animals is low.

A 10-month monitoring study was undertaken. Forty-two wells were chosen with the Stratified Random Sample (Murray, 2002) and site-location data were determined with global positioning system units (Garmin 12XL, Garmin International Inc., Kansas, USA). Water pH was measured in the field with Hanna Instruments® HI9025, whereas electrical conductivity (EC) and temperature were measured with Hanna Instruments® HI9835. The sampling periods were defined in accordance with the precipitation regime and variations in the hydrologic levels in the wells. Based on these criteria, four sampling seasons were established (March, June, September, and December).

Water samples were analyzed for total coliforms (TC), fecal coliforms (FC), and fecal streptococci (FS). Although TC is widespread in the environment, it was included in order to meet the Chilean standard requirement (NCh 409. Of 70). Aseptic sample collections were taken in sterilized flasks. Samples were held at 5 °C after being collected and for no more than 6 h until reaching the laboratory. Results were expressed in colony forming units (CFU) per 100 mL. TC, FC, and FS concentrations were analyzed with a membrane filtration technique following standard methods (Clesceri *et al.*, 1998). Aliquots (100, 10, and 1 mL) of each water sample were filtered through a 0.45 µ Millipore membrane filter. All samples were tested in triplicate. Results were reported as CFU 100

mL⁻¹. Samples that were overgrown were considered to contain > 1000 CFU 100 mL⁻¹. Colonies forming a green metallic sheen were counted as TC on m-Endo agar (Difco®, Detroit, MI, USA). To count FC, filters were placed on Petri dishes containing m-FC agar (Difco®, Detroit, MI, USA) which gave the selected colonies a blue color, whereas the selective FS count was carried out by incubating the filters in m-Enterococcus agar (Difco®, Detroit, MI, USA). Water sample analyses were performed in the microbiology laboratory of the Centro de Ciencias Ambientales (EULA) at the Universidad de Concepción.

Count data analyses were performed with STATISTICA™ StatSoft 6.0. The median was used rather than the mean to analyze the microbiological data because it basically eliminates extreme values (Smith *et al.*, 1996).

Results for TC, FC, and FS obtained in the four seasons were analyzed by looking for spatial correlations with spatial S-PLUS software using Geary's and Moran's Index (Cai and Wang, 2006). Statistical analyses were conducted to determine the relationship between bacterial concentrations and pH, electrical conductivity, temperature, and factors expected affecting concentrations or associated with the presence of indicator bacteria. These variables were treated as binomial categorical data. To further the analysis, the variables were transformed from continuous to categorical. Data included different land use activities (prairie, gardens, orchards, bare soil) within the proximity of the monitoring well (*ca.* 10 m radius); well condition (good, average, and poor); well location (highlands or lowlands); well cover (wood or cement); border height (to 15, 50, and 100 cm); casing (cement or brick); slope (to 15%, between 15% and 60%); latrine characteristics (location uphill or downhill from the well, casing); animal access in the vicinity of the well; type of animal (horses, pigs, sheep, poultry, cattle, dogs), and well-to-latrine distance (to 30 m, to 80 m). Parameters such as soil and geology were assumed to be constant because of the small differences detected at each sample site. Data were analyzed statistically by nonparametric Mann-Whitney rank-sum and Kolmogorov-Smirnov tests (Rohatgi, 1984) to determine significant differences in mean concentrations and indicator distribution found in

well groups presenting specific characteristics. Factors were ordered dichotomously. Rainfall data were collected as an additional factor likely to exert an influence on microbiological quality. The environmental variables were selected because of their expected impact on the numbers of microorganisms detected in the samples.

RESULTS AND DISCUSSION

Groundwater indicator bacteria concentrations exceeded Chilean water quality regulations in all samples (NCh 409 Of. 70). The three indicators had a detection rate of 100%, finding at least 1 CFU 100 mL⁻¹ in all tested samples. These concentrations indicated degraded groundwater quality. The existence of both FC and FS provided strong evidence of fecal contamination (Atherholt *et al.*, 2003). The presence of indicators in all four sampling seasons denoted frequent, if not continuous, fecal contamination in the ESJ watershed. There seemed to be a permanent source of fecal bacteria regularly entering the wells. Microbial data (Table 1) revealed marked variations throughout the year.

The most frequent indicator was TC. Seasonal variations in the microbial quality of water were evident, with peaks in winter for TC, FC, and FS. Variations in FC were less dramatic than in FS. Median concentrations of TC, FC, and FS increased in June (as compared to March), decreased in September, and increased again in December (Figure 2). This last increase can be attributed to higher demands on the wells during the later part of the year, combined with minimal water yields. Environmental persistence or growth of bacterial indicators during the summer months could confound the interpretation of baseline dynamics (Shanks *et al.*, 2006).

The wells exhibited a high proportion of low counts and a small number of very high counts that exerted a significant influence on the median. Indeed, bacterial indicators from natural sources do not usually occur in elevated concentrations since they come from disperse sources such as waste of warm-blooded animals (Ortiz, 2004). During transport and after retention in the soil, microorganisms are affected by environmental conditions

Table 1. Median and range of indicator bacteria concentrations in the four sampled months (CFU 100 mL⁻¹).

Sampled month	N° samples	Total coliforms		Bacterial indicator Fecal coliforms		Fecal streptococci	
		Median	Range	Median	Range	Median	Range
March	41	257	16 – 4.71×10 ³	27	1 – 1.16×10 ³	196	9 – 1.12×10 ³
June	41	501	14 – 5.00×10 ³	190	1 – 5.80×10 ³	290	20 – 1.17×10 ³
September	42	255	11 – 1.06×10 ⁴	10	1 – 3.00×10 ²	67	9 – 1.28×10 ³
December	39	440	22 – 3.60×10 ³	53	1 – 1.38×10 ³	120	5 – 1.10×10 ³

such as nutrient availability and predation (Pachepsky *et al.*, 2006). Moreover, traditional monitoring and research programs quantify the microorganism concentrations in samples using standard methods. These methods are designed to target public health and do not completely measure either clumped organisms or those associated with particles, and may not fully specify organism concentrations (Borst and Selvakumar, 2003).

Indicator concentration data fit a logistic distribution, showing a parallel evolution in the distribution of FC and FS (Figures 3, 4, 5). A descriptive criterion was chosen for this distribution.

Statistical analyses showed that FC was better correlated with TC in March, June, and December, and with FS in September. In June (winter), the three indicators showed the highest correlation. FC and TC were highly correlated. Correlation analyses revealed a strong, significant, and positive correlation between TC and FC in June (Table 2). A strong relationship between two indicators may provide some evidence that both indicators originate from the same or similar contamination sources (Francy *et al.*, 2000). Correlations between indicators, without considering the season, were very low ($r = 0.35$

between TC and FC, $r = 0.34$ between FC and FS, and $r = 0.21$ between TC and FS, $p < 0.05$). Strong correlations between indicators were obtained only when the analyses considered the season. Analysis of the annual pattern showed almost no correlations. This confirmed the importance of carrying out seasonal analyses.

By comparing indicator medians in different seasons (Kruskal-Wallis test for comparing medians), it was possible to obtain results for FC ($p\text{-value} = 2.95 \times 10^9$) which infer that seasonal medians were not equal, although FC did not change drastically with the seasons. There were differences (with a significance level of 5%) between the medians of: March/June, March/September, June/September, and September/December. FS had a $p\text{-value} = 7.95 \times 10^7$. Differences had the same significance level between the medians of: March/September, June/September, and June/December. The significant differences observed between the median concentrations of June with respect to September and December for FC and FS showed a temporal change. Median concentrations of TC did not differ significantly between seasons. Persistence of bacteria in the aquatic environment depends on various parameters, especially on the existing nutrients and temperatures (Leclerc *et al.*, 2002). The prevalence of FS, which die off more rapidly in the environment than other bacterial indicators, shows either relatively recent contamination of a source by fecal

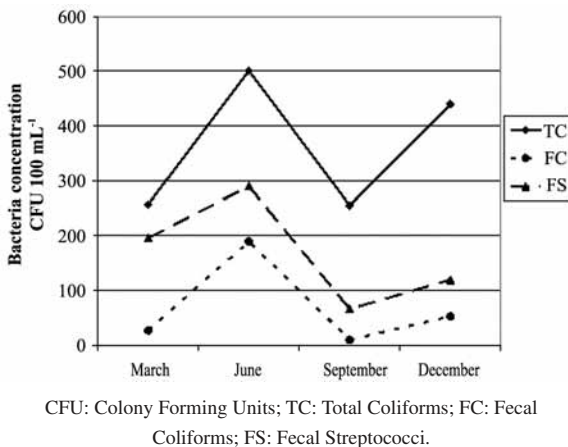


Figure 2. Median concentrations of indicator bacteria.

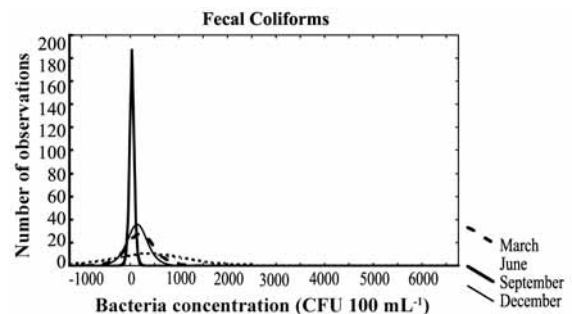


Figure 4. Logistic distribution of FC (Fecal Coliforms).

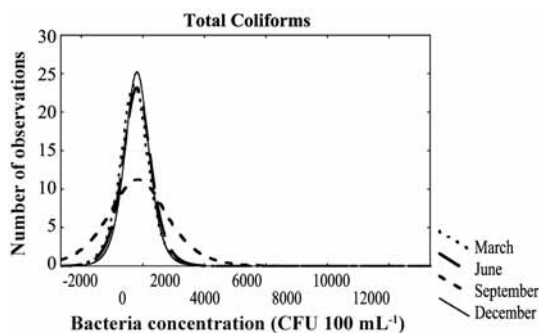


Figure 3. Logistic distribution of TC (Total Coliforms).

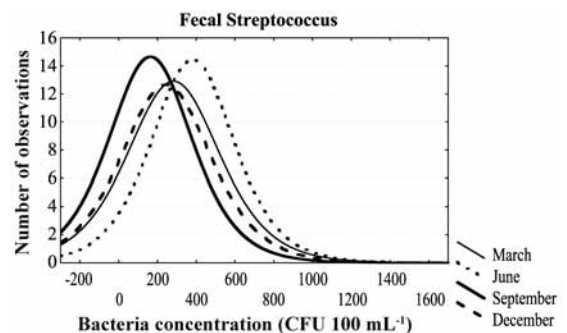


Figure 5. Logistic distribution of FS (Fecal Streptococci).

Table 2. Correlation coefficients (r) between different indicator organism concentrations.

Organism	Sample month											
	March			June			September			December		
	TC	FC	FS	TC	FC	FS	TC	FC	FS	TC	FC	FS
TC	1	0.26	0.28	1	0.91*	0.60*	1	0.23	0.41**	1	0.40**	-0.06
FC		1	0.11		1	0.53*		1	0.24		1	0.12
FS			1			1			1			1

* $p < 0.001$. ** $p < 0.05$. TC: Total coliforms. FC: Fecal coliforms. FS: Fecal streptococci.

material or a very high level of contamination possibly associated with organic matter (Conboy and Goss, 2001); the latter could have been the case in September. FC was more persistent in freshwater than FS (Anderson *et al.*, 2005). Nevertheless, in an experiment of some treatments in simulated groundwater environments by Conboy and Goss (2001), FS was able to survive for over 140 d.

Concentrations of TC, FC, and FS were not correlated with well temperature, conductivity, and pH ($p < 0.001$). Rainfall measured over the sampling period was 23.6 mm until March, 447.7 mm between March and June, 302.5 mm between June and September, and 59.9 mm between September and December (Figure 6).

The highest rainfall was recorded between May and July. FC and FS median concentrations varied over time and showed a pattern similar to that of rainfall. However, FS were more affected by rainfall than FC, although the variation patterns of FC were highly influenced by two extreme concentrations. Correlation coefficients between indicators and rainfall showed a significant relationship with FC ($r = 0.84$) and FS ($r = 0.81$). This relationship was weak for TC ($r = 0.23$) and not coupled with other factors. The high temporal variance in the collected data means that precipitation can exert an influence by providing transport energy for the potential sources. The median demonstrated that microbial water quality changes following a rainfall runoff pattern for microbial source inputs, with a marked annual cycle (Figure 6). Results revealed a strong association between bacterial concentrations in groundwater wells and rainfall through elevated concentrations in samples taken after precipitation. It can be assumed that the higher concentrations recorded in June are partly attributable to the fact that it is the wettest month of the year. These correlations suggest that bacteria were largely associated with suspended particulate materials and transported by runoff, since some coliforms in runoff are associated with particles (George *et al.*, 2004). Characteristics of the initial fecal material deposition site on the soil surface influence the infiltration, runoff, and retention rate of the microorganisms in the feces (Ferguson *et al.*, 2003). Soil surrounding wells was eroded at almost all the sites,

thereby preventing interaction between bacteria that could be transported by runoff and allowing them to eventually reach the well.

Moreover, no spatial correlations were found according to Geary's and Moran's Index. Neighboring wells were hydrologically independent. Spatial variability in the concentrations of TC, FC, and FS was not significant (Kolmogorov-Smirnov test, $p < 0.05$) between sampling sites in the highlands and lowlands of the watershed. Fecal contamination due to surface runoff implied that the phenomenon is highly responsive to rainfall intensity and duration, and will display a high degree of temporal variability. The fact that there is no significant difference between concentrations of indicators in highlands and lowlands suggests that local runoff produced the contamination rather than a landscape level phenomenon.

The analysis of the relationship between bacterial indicator levels and environmental characteristics presents several statistical challenges. Due to the complex nature of FC destination and transport, empirical methods such as regression models are unable to build up reliable load-concentration relationships (Bai and Lung, 2006). However, factors (Table 3) were recorded which were expected to affect concentrations or be associated with the presence of indicator bacteria since these offer preliminary insight into the causes of well contamination.

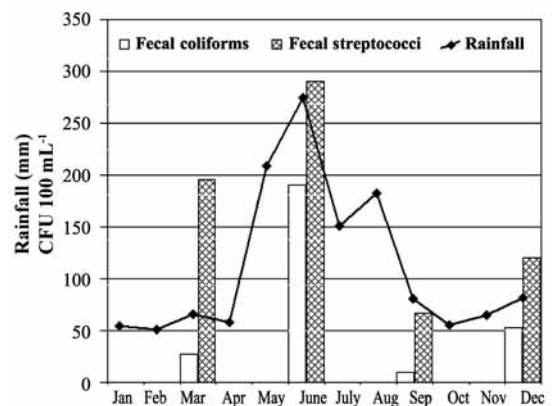


Figure 6. Rainfall and seasonal variability of indicator bacteria concentrations.

Table 3. Landscape and management factors expected to affect concentrations or be associated with the presence of indicator bacteria.

Well	Land use	Slope	AA	Animal type	Well condition	Well cap	Well cover	B h	LU	D	L casing
1	Prairie	5	Yes	H	Good	Wood	Cement	45	Yes	85	Cement
2	Orchard	15	Yes	Pi	Good	Wood	Cement	100	Yes	11	Cement
3	Garden	5	No	-	Good	Wood	Brick	30	No	42	None
4	Orchard	5	Yes	Po	Good	Cement	Cement	60	No	84	None
5	Orchard	9	No	-	Regular	Cement	Brick	30	Yes	44	Cement
6	Prairie	15	Yes	S	Regular	Wood	Cement	30	No	40	None
7	Prairie	10	Yes	S	Poor	Wood	Cement	40	No	52	None
8	Bare soil	26	No	-	Good	Cement	Cement	3	Yes	70	None
9	Orchard	10	Yes	Po	Regular	Cement	Cement	70	Yes	28	None
10	Orchard	18	No	-	Regular	Wood	Cement	75	Yes	83	None
11	Orchard	40	Yes	Po	Poor	Wood	Cement	40	Yes	62	Cement
12	Orchard	35	Yes	Po, C	Poor	Wood	Cement	40	Yes	85	None
13	Orchard	35	Yes	Po	Regular	Cement	Cement	40	Yes	50	None
14	Orchard	40	Yes	Po, pi	Good	Cement	Cement	60	Yes	62	Cement
15	Bare soil	18	No	-	Regular	Wood	Cement	50	Yes	82	None
16	Prairie	30	Yes	Po	Poor	Wood	Brick	18	Yes	120	None
17	Bare soil	45	Yes	Po, C	Good	Cement	Cement	40	No	86	None
18	Prairie	5	Yes	Po, C	Good	Cement	Cement	50	No	86	None
19	Prairie	53	Yes	Po, pi	Regular	Wood	Cement	5	Yes	60	None
20	Orchard	45	No	-	Good	Cement	Brick	15	Yes	48	None
21	Orchard	25	No	-	Good	Cement	Brick	60	Yes	43	None
22	Bare soil	5	Yes	H	Good	Cement	Cement	20	Yes	70	None
23	Orchard	0	No	-	Regular	Cement	Brick	40	No	22	None
24	Prairie	18	No	-	Regular	Cement	Brick	60	Yes	18	None
25	Prairie	15	Yes	Po, D	Good	Cement	Cement	70	Yes	50	None
26	Garden	10	Yes	Po	Good	Cement	Cement	40	No	23	Cement
27	Orchard	23	Yes	C	Good	Cement	Cement	60	Yes	62	Cement
28	Orchard	4	Yes	Po	Regular	Wood	Brick	80	Yes	79	Cement
29	Orchard	17	Yes	Po	Regular	Wood	Cement	55	Yes	80	Cement
30	Prairie	18	Yes	Po, C	Poor	Wood	Cement	10	Yes	91	None
31	Prairie	40	Yes	Po, C	Good	Wood	Cement	50	Yes	133	None
32	Orchard	22	Yes	Po, S	Regular	Wood	Cement	50	Yes	10	Cement
33	Orchard	12	Yes	Po	Good	Cement	Cement	20	Yes	42	None
34	Bare soil	13	Yes	S	Poor	Cement	Cement	5	Yes	54	None
35	Garden	20	No	-	Good	Wood	Cement	50	Yes	26	None
36	Orchard	15	Yes	D	Good	Cement	Cement	60	Yes	39	None
37	Prairie	5	No	-	Good	Cement	Cement	100	No	10	Cement
38	Orchard	18	Yes	Po	Poor	Wood	Cement	90	Yes	41	None
39	Prairie	35	No	-	Regular	Cement	Brick	120	No	45	None
40	Orchard	40	Yes	Po, D	Poor	None	Cement	45	Yes	11	None
41	Orchard	5	No	-	Poor	None	Cement	12	Yes	63	None
42	Orchard	15	No	-	Regular	Wood	Cement	30	No	82	None

AA: animal access to the well. Bh: well border height. LU: latrine uphill from the well. D: distance between well and the closest latrine. L casing: latrine casing. Animal type: H: horses, Pi: pigs, Po: poultry, S: sheep, C: cattle, D: dogs.

The distance between wells and latrines is highly variable, ranging from a minimum of 10 to 133 m. Seventy-four percent of the latrines in the sampled households had brick casing. Hence, they were not sealed. On at least one of the four sampling dates, animals were observed around approximately 67% of the wells. Table 4 demonstrates that characteristics with $p < 0.052$ were considered to be statistically significant. Statistical analysis of the data showed that five factors are likely to influence the concentration of bacteria in groundwater: animal access close to the wells (specifically pigs and poultry); land use; bricks used for well casing; latrine-to-well distance; and a slope up to 15%. Only two of these factors showed a highly significant ($p < 0.01$) association with the presence of the bacterial indicators: animal access close to the well in June and a latrine-to-well distance of < 80 m in December.

These results suggest that the most important factors affecting well vulnerability to bacterial contamination were those related to the well itself: construction and site management. In the month when the indicator concentrations are the highest, the factors potentially influencing these levels are animal access (specifically poultry) and well casing. Some wells have a brick casing instead of cement, which does not seal them sufficiently and allows water runoff from the surroundings to enter. Statistically, contamination levels were more closely tied to animal access in the vicinity of wells and the well casing material than to land use or distance between wells and latrines. Livestock grazing practices creates a diffuse source of fecal contamination to watersheds (Tian *et al.*, 2002; Harter *et al.*, 2002). Pathogens from animal feces may enter waterways by direct deposition or as a result

of overland runoff containing fecal material deposited in the watershed. The FC:FS ratio as used by (Donderski and Wilk, 2002; Troussellier *et al.*, 2004) showed that the source of indicator bacteria is mostly animal, followed by mixed sources. Considering that a great number of wells have fences to prevent animal access, wildlife cannot be disregarded as a source. Cox *et al.* (2005) showed that poultry fecal samples have a higher FC concentration (median 1.1×10^8 CFU g^{-1} wet wt) than those of other domestic animals (median for adult cattle 1.8×10^5 CFU g^{-1} wet wt, pigs 7.1×10^6 CFU g^{-1} wet wt, and sheep 6.6×10^5 CFU g^{-1} wet wt). This could explain the significance of poultry access to the wells as a factor affecting indicator counts. Furthermore, Wheeler *et al.* (2002) demonstrated that *Enterococcus faecalis* had a limited host range and was found in humans, dogs, and chickens.

Land use in the watershed also affected the extent of fecal contamination, but not as strongly as the other factors described above. A pattern did not emerge in spite of the fact that three different land uses were significant. Latrines appear to have little influence on the presence and level of bacterial indicators, suggesting that latrines can also be a potential source of microbial contamination in groundwater. Other factors not considered in this study may also affect bacterial concentrations in well water.

These data provide new information by relating indicator bacteria loads for certain factors at specific times of the year. The fact that the most significant indicator related to a factor was TC in March and in December, FC in June, and FS in September, suggests that fecal contamination is mostly a winter phenomenon.

Table 4. Factors with significant differences between the means of indicator bacteria concentrations.

Month	Indicator	Factor	P	Test
March	Fecal streptococcus	Land use: bare soil	0.038	Mann-Whitney
March	Total coliforms	Pig access	0.015	Mann-Whitney
June	Fecal coliforms	Animals close to well	< 0.005	Kolmogorov-Smirnov
June	Fecal coliforms	Animals close to well	0.021	Mann-Whitney
June	Fecal coliforms	Poultry access	< 0.05	Kolmogorov-Smirnov
June	Fecal coliforms	Poultry access	0.052	Mann-Whitney
September	Fecal streptococcus	Land use: orchard	0.051	Mann-Whitney
September	Fecal streptococcus	Well casing material (brick)	< 0.05	Kolmogorov-Smirnov
September	Fecal streptococcus	Well casing material (brick)	0.041	Mann-Whitney
September	Fecal streptococcus	Latrine-to-well distance < 80 m	0.044	Mann-Whitney
December	Fecal streptococcus	Land use: garden	0.039	Mann-Whitney
December	Fecal streptococcus	Slope $< 15\%$	0.041	Mann-Whitney
December	Total coliforms	Well casing material (brick)	< 0.05	Kolmogorov-Smirnov
December	Total coliforms	Well casing material (brick)	0.033	Mann-Whitney
December	Total coliforms	Latrine-to-well distance < 80 m	< 0.05	Kolmogorov-Smirnov
December	Total coliforms	Latrine-to-well distance < 80 m	0.007	Mann-Whitney

CONCLUSIONS

There is widespread groundwater contamination in the ESJ watershed. The microbiological quality of the sampled wells was impaired with regard to Chilean standards.

A seasonal trend was identified. Concentrations of FC and FS varied over time and showed a pattern similar to rainfall which appeared to exert a local influence on the indicator concentrations. FS were more affected by rainfall than FC.

The lack of a significant difference between wells located uphill and downhill suggests that contamination is not a result of surface runoff from upgradient areas. Our results indicate that one cause of microbial contamination in well water is manure bacteria entering directly through local surface runoff.

There was no spatial correlation between wells, showing that there were no identified groups of wells which maintained certain concentration tendencies.

The present study shows that the analysis of microbial data in combination with basic environmental and management data can provide preliminary insight into the causes of fecal contamination in groundwater. In fact, indicator counts turned out to be significantly related to certain watershed features during specific months. Inherent well site characteristics and its surroundings, as well as rainfall are the main factors that affect groundwater quality in the ESJ watershed.

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RESUMEN

Contaminación fecal en agua subterránea en una pequeña cuenca de secano rural en Chile Central. Se realizó una investigación de la calidad microbiológica de las aguas subterráneas en una cuenca rural chilena. En esta cuenca prácticamente no había otra fuente de agua disponible. En 42 pozos seleccionados al azar, se midieron niveles de bacterias indicadoras en cuatro temporadas distintas durante el año 2005. Las bacterias incluyeron coliformes totales (TC), coliformes fecales (FC) y *Escherichia coli* (FS). El objetivo fue caracterizar la calidad microbiológica del agua subterránea y relacionar los indicadores con ciertas propiedades y el manejo de la cuenca que pueden afectar la calidad del agua. La dinámica temporal de la contaminación fue determinada mediante análisis estadístico de la concentración de organismos

indicadores. Las relaciones entre indicadores bacteriales presentes en el agua de los pozos y otras variables fueron analizadas con pruebas no paramétricas. En todas las muestras se detectaron TC, FC y FS, indicando que los pozos han estado contaminados con material fecal de humanos y animales. La distribución de frecuencia de los microorganismos se ajustó a una distribución logística. Las concentraciones muestran una base temporal con niveles variables entre temporadas, con una mayor concentración en invierno. La causa de la contaminación se puede asociar al fácil acceso de los animales domésticos a los pozos, y a su material de revestimiento permeable. La escorrentía local de las precipitaciones mostró tener una influencia directa sobre la concentración de los microorganismos en los pozos y en la concentración de los indicadores bacteriales encontrados en los pozos.

Palabras clave: contaminación biológica, bacteria, calidad del agua, contaminación ambiental.

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PHOSPHATE FERTILIZATION CAN INCREASE YIELD OF PRODUCTIVE GRASS PEA (*Lathyrus sativus* L.) CROPS IN P-RETENTIVE SOILS

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ABSTRACT

The effect of P fertilization on grass pea (*Lathyrus sativus* L.) yield and yield components was evaluated on soils with low P availability and high P retention capacity in small-scale farms of the Araucanía Region in southern Chile. Trials were conducted during 2000-2001, 2001-2002, and 2002-2003, in six sites; three sites in Lumaco and three in the Selva Oscura area. Six rates of P (0, 21.8, 43.6, 65.4, 87.2, and 109.0 kg ha⁻¹) were evaluated in a randomized complete block design with four replicates. Grass pea cv. Luanco-INIA was sown at 47 seeds m⁻². Mean grain yield for all trials was 2456 kg ha⁻¹. Phosphate fertilization increased grass pea grain yield in both areas during 2000 and 2001. There was no significant effect in 2002. The 2002 cropping season had an unusually high spring–summer rainfall, which may have enhanced the P mineralization rate from organic soil fraction, and thus P availability. According to this study, grass pea crops in soils with < 10 mg kg⁻¹ of available P-Olsen **should respond** to P fertilization.

Key words: grass pea, *Lathyrus*, phosphorus, neglected crops, cool-season legumes.

INTRODUCTION

Grass pea is a grain legume crop used for human and animal consumption since ancient times (Hanbury *et al.*, 2000). The presence of β -N-oxalyl-L- α , β -diaminopropionic acid (β -ODAP) in grass pea seeds is thought to increase vulnerability to neurolethyrism, a neurodegenerative disease (Lambein *et al.*, 2007). As a result, lowODAP lines (Campbell *et al.*, 1994) and cultivars (Siddique *et al.*, 2006) have been found that may enhance interest in this protein crop. The potential of grass pea for animal feed also depends on the achievement of yields that make it competitive with other protein-rich ingredients.

Grass pea is grown on a wide range of soils, including those with low fertility and poor structure (Siddique *et al.*, 1996). In southern Chile, grass pea is cultivated by small farmers with limited resources, on typically eroded soils as a result of poor management. The Araucanía Region (37°30'–39°30' S) has a particularly high number of such small farmers and its soils are characterized by low levels of available P-Olsen (Montenegro, 1991) though relatively high grain yields are often achieved. Hence, P fertilization is a determining factor in the yield of most crops.

Information on grass pea response to P fertilization is very scarce. Its effect on grain yield and yield

components is largely unknown, particularly in soils with high P retention capacity. The comprehensive review by Campbell (1997) does not refer to P fertilization, but emphasizes that grass pea is considered a hardy crop requiring low or zero inputs. However, Sarkar *et al.* (2003) found that applying P increased grass pea grain yield grown in an Entisol of India with pH 7.5, 0.53% organic carbon, and 26 kg ha⁻¹ of P₂O₅. In Chile, a clear effect of up to 65.4 kg ha⁻¹ P on grass pea grain yield was reported by Ellena (1983) in an Andisol of Valdivia. Later, Montenegro *et al.* (2001) explained in a preliminary report that P fertilization was associated to higher grass pea yields in Araucanía soils with high P retention capacity. Krarup (2002) did not find any grass pea response to P fertilization in a soil of the Valdivia series belonging to the medial, mesic of the Duric Hapludands family (CIREN, 2003) with P-Olsen availability of 12 mg kg⁻¹ or greater.

Araucanía Region soils have high levels of P retention, generally above 70% (Sadzawka *et al.*, 1999) reaching levels as high as 95% in areas such as Victoria. Phosphorus retention of 70-99% has been reported for the A soil horizon in Andisols of southern Chile (Pino *et al.*, 1998; Besoain and Sadzawka, 1999). A wide range of farmers was surveyed in the Araucanía Region indicating that 90% of soils have available P-Olsen < 15 mg kg⁻¹ (Sadzawka *et al.*, 1999). Moreover, 60% of such soils have P-Olsen < 10 mg kg⁻¹ which is considered a critical level for most crops. This level is frequent in small farms, with a mean of 7.5 mg kg⁻¹ for grass pea producers in

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Lumaco (A. Montenegro, unpublished data, 2000), one of the areas of the Araucanía Region where the reported experiments were conducted.

Phosphorus is generally absorbed by crop plants to a moderate extent compared to other macronutrients. However, in P-retentive soils such as those in southern Chile, P fertilizers should be applied at high rates, for example, up to 87 kg ha⁻¹ P for wheat (Montenegro *et al.*, 1999a) and oilseed rape (Montenegro *et al.*, 1999b). They should be localized in the furrow to increase efficiency. Consequently, it is important to gather information on the magnitude of yield response of grass pea to P fertilization in soils where its availability is limited.

MATERIALS AND METHODS

The study was conducted under dryland conditions during three cropping seasons (2000-2001, 2001-2002, and 2002-2003 referred hereafter as 2000, 2001, and 2002) in Lumaco (38°08'S, 72°55'W) and Selva Oscura (38°21'S, 72°11'W), Araucanía Region, located about 150 km NW and 60 km NE of Temuco, respectively. Experiments in Lumaco were conducted on a silty clay loam Inceptisols, Lumaco Series, belonging to the fine, mixed, termic, mesic of the Fluventic Dystrudepts family (CIREN, 2002). Experiments in Selva Oscura were conducted on a silty clay loam Andisols, Victoria Series, belonging to the medial, mesic of the Typic Durudands family (CIREN, 2002). Methodology for chemical determination was in

accordance with Sadzawka *et al.* (2006). Specific sites within Lumaco and Selva Oscura were different each year.

The chemical characterization of soils (0-20 cm depth) is shown in Table 1. Available P-Olsen in Lumaco soils was low in all seasons. In Selva Oscura, available P-Olsen was considered low to medium in 2000, and low in 2001 as well as 2002. Traditional crops, like wheat (*Triticum aestivum* L.) and oilseed rape (*Brassica napus* L.), for which more agronomic information is available, are expected to clearly respond to P fertilization in soils at these P-Olsen levels. In Lumaco, exchangeable K was low in 2000 and 2002, medium in 2001, and extractable S was low, whereas Al saturation was low in 2000 and 2001, but high in 2002. In Selva Oscura, soils had relatively high exchangeable K, low extractable S, and medium to high Al saturation.

Luanco-INIA, a large-seeded grass pea cultivar (Mera *et al.*, 2003), was sown at 47 seeds m⁻². Seeds were treated with carboxin + thiram, fipronil, and inoculated just prior to sowing with a cocktail of Rhizobium strains isolated from grass pea nodules provided by the Universidad de Concepción, Chillán, Chile. Sowing dates in Lumaco were 17 May 2000, 16 May 2001, and 7 May 2002. Sowing dates in Selva Oscura were 11 August 2000, 7 August 2001, and 3 September 2002. A randomized complete block design was used with four replicates, except in Lumaco in 2000 and 2001 where only three replicates were carried out. Plots had six rows, 4 m long and spaced

Table 1. Chemical characterization of soils (0-20 cm depth) from three different sites in two areas of southern Chile where experiments of P fertilization in grass pea were conducted during 3 years.

Chemical parameters and its unit	Lumaco			Selva Oscura		
	2000	2001	2002	2000	2001	2002
P-Olsen, mg kg ⁻¹	9	5	5	11	6	4
Organic C, %	2.9	2.3	3.5	7.0	6.4	4.1
pH (water)	5.8	5.8	5.2	5.5	5.8	5.6
Exchangeable Ca, cmol kg ⁻¹	4.9	3.35	1.97	4.9	5.18	9.05
Exchangeable Mg, cmol kg ⁻¹	1.04	1.3	0.76	1.45	0.8	2.16
Exchangeable Na, cmol kg ⁻¹	0.26	0.08	0.15	0.11	0.07	0.13
Exchangeable K, cmol kg ⁻¹	0.07	0.52	0.13	0.88	0.96	0.38
Sum of bases, cmol kg ⁻¹	6.3	5.25	3.01	7.38	7.01	11.72
Exchangeable Al, cmol kg ⁻¹	0.09	0.07	1.24	0.79	0.28	1.2
Effective CEC, cmol kg ⁻¹	6.4	5.32	4.25	8.17	7.29	12.92
Al saturation, %	1.4	1.3	29.2	9.7	3.8	9.3
Extractable S, mg kg ⁻¹	2.4	6.5	-	4.5	1.3	-
Available Zn, mg kg ⁻¹	0.3	0.1	-	1.5	0.1	-
Available B, mg kg ⁻¹	0.4	0.2	-	0.5	0.2	-
Available Cu, mg kg ⁻¹	3	1.4	-	2.5	0.6	-

Effective CEC: effective cation exchange capacity; Aluminium saturation percentage is Exchangeable Al divided by Effective CEC multiplied by 100.

35 cm apart. Treatments were 0, 21.8, 43.6, 65.4, 87.2, and 109.0 kg ha⁻¹ P, as banded superphosphate below the seed (46% P₂O₅). Base fertilization was 83 kg ha⁻¹ K and 36 kg ha⁻¹ S as broadcast K₂SO₄ in all experiments and for all P treatments. In 2002, AI soil saturation recommended a liming treatment, so 3.5 and 3.0 t ha⁻¹ CaCO₃ with an agronomic value of 92.44% was applied in Lumaco and Selva Oscura, respectively.

Aboveground biomass samples were taken in Lumaco during grass pea flowering to estimate nutrient contents only in year 2000. Grain was harvested 17 January 2001, 2 January 2002, and 16 January 2003 in Lumaco, and 26 January 2001, 21 January 2002, and 25 February 2003 in Selva Oscura. Nutrient content in biomass and grain was estimated at both locations only in 2000. Nitrogen content in biomass and grain was determined by digestion with sulphuric acid and Kjeldahl. Phosphorus content in aboveground biomass and grain was determined by calcination, digestion with HCl, and colorimetry by vanadate. Potassium, Ca, Mg, Zn, and Cu in aboveground biomass and grain were determined by calcination, digestion with HCl, atomic absorption, and emission spectrophotometry. Boron in the aboveground biomass and the grain was determined by calcination, digestion with HCl, and colorimetry with azomethine-H (Sadzawka *et al.*, 2007). Samples of aboveground biomass were taken in Lumaco and Selva Oscura in 2001 during flowering in order to estimate aboveground dry biomass per hectare. Grain yield was estimated using 3.5 m of the four central rows at 14% moisture. Yield components were estimated from a row random sample of 25 cm. Nutrient content of

grains from both areas was determined in 2000. Analysis of variance and regression were performed with SAS (SAS Institute, 1992).

RESULTS AND DISCUSSION

In 2000 and 2001, P fertilization increased grain yields in both areas, in agreement with previous findings (Ellena, 1983). However, no significant effect was found at any site in 2002 (Tables 2 and 3).

Grain yield mean was considerably higher in Lumaco (2908 kg ha⁻¹) than Selva Oscura (1268 kg ha⁻¹) in 2000. The higher level of AI saturation in Selva Oscura (Table 1) may have been detrimental to grain yield. Furthermore, soil structure in Selva Oscura was altered by extremely high rainfall in June 2000 (Figure 1), and a superficial soil crust resulted from the impact of raindrops (Casanova *et al.*, 2006), probably affecting root development. In 2000, the effect of increasing P fertilization rates on grain yield fitted a quadratic model response with Equation [1] for Lumaco and Equation [2] for Selva Oscura:

$$y = 2335.39 + 18.2995 P - 0.097557 P^2, \text{ with } R^2 = 0.60 \quad [1]$$

$$y = 875.14 + 8.300078 P - 0.01356 P^2, \text{ with } R^2 = 0.77 \quad [2]$$

where y is the grain yield (kg ha⁻¹), P is the amount of P applied (kg P ha⁻¹), and R^2 is the coefficient of determination for the equation.

Contrary to 2000, there were higher mean yields in 2001 in Selva Oscura (2465 kg ha⁻¹) than in Lumaco (1687 kg ha⁻¹). Water availability in 2001 was less than

Table 2. Effect of P fertilization on stand, grain yield, and grain weight of grass pea cv. Luanco-INIA in Lumaco, southern Chile, during 3 years.

Applied phosphorus	2000			2001			2002		
	Stand plants	Grain yield	Grain weight	Stand plants	Grain yield	Grain weight	Stand plants	Grain yield	Grain weight
kg ha ⁻¹ P	plant m ⁻²	kg ha ⁻¹	mg	plant m ⁻²	kg ha ⁻¹	mg	plant m ⁻²	kg ha ⁻¹	mg
0.0	41.4	2359	327	32.8	1036	286	40.2	3248	285
21.8	40.6	2695	333	33.1	1553	284	38.2	3126	305
43.6	40.8	2841	340	29.2	1553	279	38.6	3598	288
65.4	39.7	3155	341	32.0	1700	281	40.8	3482	283
87.2	38.8	3282	346	30.4	1945	294	38.6	3425	297
109.0	39.5	3114	330	35.0	2334	296	34.5	3014	292
Mean	40.1	2908	336	32.1	1687	287	38.5	3316	292
CV, %	4.7	7.8	2.7	12.3	16.4	6.3	12.40	14.0	6.1
F	0.79	6.91	2.04	1.10	9.83	0.59	0.86	0.93	0.88
P > F	ns	0.01	ns	ns	< 0.01	ns	ns	ns	ns

Grain weight is a mean value calculated from a random sample of 500 grains. ns = non significant.

CV: coefficient of variation. F: ratio of treatments mean square and experimental error mean square.

Table 3. Effect of P fertilization on stand, grain yield, and grain weight of grass pea cv. Luanco-INIA in Selva Oscura, southern Chile, during 3 years.

Applied phosphorus	2000			2001			2002		
	Stand plants	Grain yield	grain weight	Stand plants	Grain yield	grain weight	Stand plants	Grain yield	grain weight
kg ha ⁻¹ P	plant m ⁻²	kg ha ⁻¹	mg	plant m ⁻²	kg ha ⁻¹	mg	plant m ⁻²	kg ha ⁻¹	mg
0.0	38.9	870	354	44.1	2143	298	45.7	2796	368
21.8	40.9	1065	365	39.2	2126	292	45.6	2952	357
43.6	37.7	1203	370	44.9	2386	297	42.3	3490	353
65.4	35.5	1342	365	35.9	2535	309	40.7	2833	359
87.2	38.2	1521	356	45.3	2966	299	43.6	3070	369
109.0	39.8	1610	370	43.6	2634	307	45.7	3584	371
Mean	38.5	1268	363	42.2	2465	300	43.9	3095	363
CV, %	5.1	11.9	2.8	11.0	9.0	3.2	11.1	13.9	5.6
F	3.61	13.68	1.8	2.65	8.26	1.69	0.74	1.98	0.55
P > F	0.05	0.01	ns	ns	< 0.01	ns	ns	ns	ns

Grain weight is a mean value calculated from a random sample of 500 grains. ns = non significant. CV: coefficient of variation. F: ratio of treatments mean square and experimental error mean square.

in 2000, particularly in Lumaco, which probably caused the lower mean grain yield there in 2001, as compared with 2000 and 2002 (Table 2). The effect of increasing P fertilization rates on grain yield in Lumaco in 2001 fitted a linear model response and a quadratic model response for Selva Oscura with Equations [3] and [4], respectively:

$$y = 1129.10476 + 10.23309 P, \text{ with } R^2 = 0.63 \quad [3]$$

$$y = 1890.05955 + 15.48615 P - 0.06904 P^2, \quad [4] \\ \text{with } R^2 = 0.54$$

where y, P, and R² are described Equations [1] and [2].

There was no significant effect of P fertilization on grain yield at any site. Unlike previous years, liming was applied in 2002 due to the high percentage of Al saturation found at both sites (Table 1). As a result, plant growth at both sites was more vigorous during 2002 than in previous years. Although not the aim of the present experiment, development of the grass pea plant was found to be severely limited by medium to high Al saturation and its associated high soil acidity. As a consequence, corrective liming was very effective. In addition, rainfall in October-December 2002 was unusually high (Figure 1) and soil moisture was still abundant during the favorable temperatures of late spring. These conditions may have enhanced P mineralization from organic soil fraction. Thus, the lack of response to added P during 2002 may have resulted from more available P in the soil. This would explain the very high grain yields (3248 and 2796 kg ha⁻¹ in Lumaco and Selva Oscura, respectively) achieved by controls where P was not applied.

Phosphate fertilization did not have a significant effect on grain weight in any site or year despite the relatively

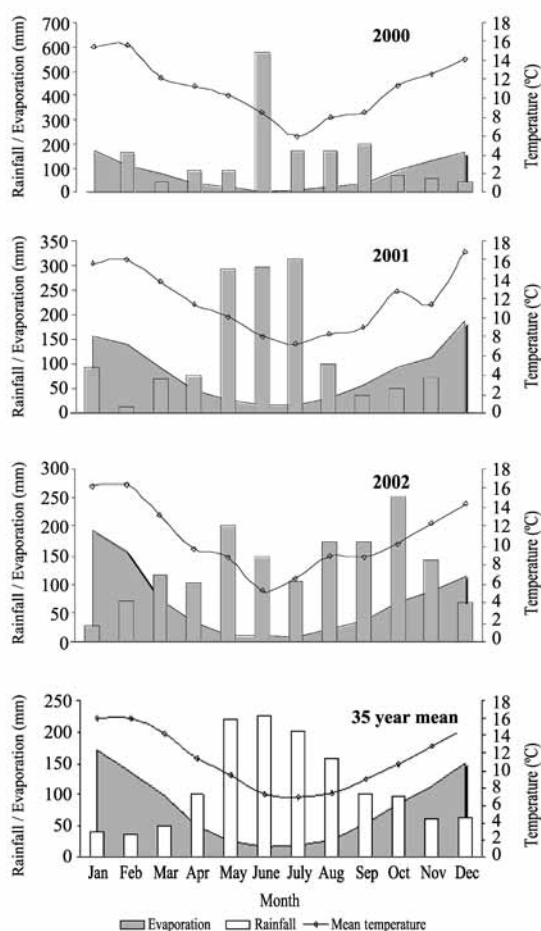


Figure 1. Monthly rainfall, evaporation, and mean temperature during 2000, 2001, 2002, and 35-yr mean at Carillanca, La Araucanía Region, southern Chile.

Table 4. Nutrient content of aboveground biomass of grass pea cv. Luanco-INIA during flowering stage in Lumaco, southern Chile in 2000-2001 for six rates of P fertilization.

Applied phosphorus	N	P	K	Ca	Mg	Zn	Mn	Cu	B
kg ha ⁻¹ P	%			mg kg ⁻¹					
0.0	2.18	0.13	0.54	0.51	0.27	13.0	67.0	5.0	31.0
21.8	1.83	0.10	0.51	0.48	0.26	12.0	65.0	5.0	32.0
43.6	2.31	0.09	0.53	0.62	0.25	12.0	83.0	4.0	34.0
65.4	2.17	0.09	0.51	0.53	0.25	12.0	74.0	4.0	31.0
87.2	1.98	0.10	0.46	0.58	0.30	14.0	73.0	4.0	35.0
109.0	2.18	0.11	0.47	0.56	0.27	13.0	67.0	4.0	32.0
Mean	2.11	0.10	0.50	0.55	0.27	12.7	71.5	4.3	32.5

Data are from a compound sample of three replicates.

Table 5. Nutrient accumulation by aboveground biomass of grass pea cv. Luanco-INIA up to flowering stage in Lumaco, southern Chile in 2000-2001 for six rates of P fertilization.

Applied phosphorus	N	P	K	Ca	Mg	Zn	Mn	Cu	B
kg ha ⁻¹ P	g m ⁻²			mg m ⁻²					
0.0	14.9	0.89	3.7	3.5	1.8	8.9	45.6	3.4	21.1
21.8	17.0	0.93	4.7	4.5	2.4	11.1	60.3	4.6	29.7
43.6	22.7	0.88	5.2	6.1	2.5	11.8	81.4	3.9	33.4
65.4	20.3	0.84	4.8	5.0	2.3	11.2	69.3	3.7	29.0
87.2	25.6	1.30	6.0	7.5	3.9	18.1	94.5	5.2	45.3
109.0	23.5	1.20	5.1	6.0	2.9	14.0	72.3	4.3	34.5
Mean	20.7	1.00	4.9	5.4	2.6	12.5	70.6	4.2	32.2

Nutrient accumulation was calculated by multiplying nutrient content by aboveground biomass.

low coefficients of variation for this trait. The 3-yr mean grain weight was higher in Selva Oscura (342 mg) than in Lumaco (305 mg), due to the relatively higher water availability during the pod filling stage in Selva Oscura. Variations in grain yield were not related to mean grain weight nor associated to the number of grains per pod, which had a mean of ~1.6. The number of pods per plant was the yield component that largely explained variations in grain yield. In 2000, plots that yielded 900-1600 kg ha⁻¹ had 6-12 pods per plant, whereas those yielding 2300-3300 kg ha⁻¹ had 13-24 pods per plant (data not shown). This noticeable difference was observed in plots with similar stands of 38-40 plants m⁻².

Phosphate fertilization did not affect aboveground biomass in 2000, but did in 2002. In Lumaco and Selva Oscura, dry matter at flowering increased from 2919 kg ha⁻¹ in controls where P was not applied to 7483 kg ha⁻¹ with 109 kg P ha⁻¹ and 3489 to 5560 kg ha⁻¹, respectively (data not shown). Plant height increased at least 10 cm as a result of P fertilization. Controls with no P were 47 cm tall during full flowering, whereas plots receiving P measured 57-63 cm with no significant differences between P rates.

The chemical characterization of the aboveground biomass during grass pea flowering in Lumaco in 2000 is shown in Table 4. Phosphate fertilization did not apparently affect neither N content, which was relatively high nor P, K, Ca, Mg, and micronutrients. Nutrient absorption by the aboveground biomass up to the flowering stage in Lumaco is presented in Table 5. In general, macro and micronutrients appeared to be absorbed to a greater extent with higher rates of P fertilization.

Nutrient content in the grass pea grain from both sites was unaffected by P treatments in 2000 (Table 6). However, mean N content of grain from Selva Oscura was less than that from Lumaco, probably due to limitations on symbiotic N fixation from high soil acidity associated with the above-mentioned Al saturation condition. On the contrary, P, K, Zn, and Cu were notably greater in grain from Selva Oscura than from Lumaco, and this was true to a lesser extent for Ca, Mg, Mn, and B. In general, nutrient absorption by grass pea grain was higher with increased P fertilization (Table 7), due to the better yield associated with it. Absorption means were lower in Selva Oscura because of lower yields.

Table 6. Nutrient content of grass pea grain cv. Luanco-INIA in Lumaco and Selva Oscura, southern Chile in 2000-2001 for six rates of P fertilization.

Applied phosphorus	N	P	K	Ca	Mg	Zn	Mn	Cu	B
kg ha ⁻¹ P	%						mg kg ⁻¹		
Lumaco									
0.0	4.0	0.23	0.67	0.10	0.09	23	22	5	6
21.8	4.2	0.28	0.73	0.10	0.09	24	19	5	8
43.6	3.9	0.20	0.67	0.11	0.09	19	22	5	8
65.4	4.5	0.20	0.69	0.10	0.08	21	22	5	9
87.2	4.0	0.23	0.65	0.10	0.09	24	21	5	8
109.0	4.2	0.18	0.67	0.11	0.09	20	23	5	8
Mean	4.1	0.22	0.68	0.10	0.09	21.8	21.5	5.0	7.8
Selva Oscura									
0.0	3.6	0.37	0.90	0.13	0.11	32	24	8	10
21.8	3.6	0.37	0.90	0.12	0.11	31	23	7	8
43.6	3.3	0.36	0.91	0.12	0.11	30	24	8	9
65.4	3.6	0.37	0.91	0.11	0.11	30	24	7	9
87.2	3.8	0.34	0.91	0.12	0.11	30	22	7	11
109.0	3.8	0.36	0.90	0.11	0.11	30	25	7	10
Mean	3.6	0.36	0.91	0.12	0.11	30.5	23.7	7.3	9.5

Data are from a compound sample of three (Lumaco) and four (Selva Oscura) replicates.

Table 7. Nutrient accumulation by grass pea grain cv. Luanco-INIA in Lumaco and Selva Oscura, southern Chile, 2000-2001, for six rates of P fertilization.

Applied phosphorus	N	P	K	Ca	Mg	Zn	Mn	Cu	B
kg ha ⁻¹ P	kg ha ⁻¹						g ha ⁻¹		
Lumaco									
0.0	81	4.7	13.6	2.0	1.8	47	45	10	12
21.8	97	6.5	16.9	2.3	2.1	56	44	12	19
43.6	95	4.9	16.4	2.7	2.2	46	54	12	20
65.4	122	5.4	18.7	2.7	2.2	57	60	14	24
87.2	113	6.5	18.3	2.8	2.5	68	59	14	23
109.0	113	4.8	17.9	2.9	2.4	54	62	13	21
Mean	103.5	5.5	17.0	2.6	2.2	54.7	54.0	12.5	19.8
Selva Oscura									
0.0	27	2.8	6.7	1.0	0.8	24	18	6	8
21.8	33	3.4	8.2	1.1	1.0	29	21	6	7
43.6	34	3.7	9.4	1.2	1.1	31	25	8	9
65.4	42	4.3	10.5	1.3	1.3	35	28	8	10
87.2	50	4.4	11.9	1.6	1.4	39	29	9	14
109.0	53	5.0	12.5	1.5	1.5	42	35	10	14
Mean	39.8	3.9	9.9	1.3	1.2	33.3	26.0	7.8	10.3

Nutrient accumulation was calculated by multiplying nutrient content by grain yield.

CONCLUSIONS

Phosphate fertilization was associated with higher grass pea yields in soils with low P availability and retention capacity. The grain yield potential of grass pea in southern Chile was confirmed to be quite high, although grain yield was linked to the amount of rainfall during the cropping season. In a dryland area, high rainfall in spring may decrease the effect of P fertilization on grain yield. With a limited water supply, a growing condition that is frequent for this crop, grass pea crops should respond to P fertilization in soils with available P-Olsen $\leq 10 \text{ mg kg}^{-1}$.

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RESUMEN

La fertilización fosfatada puede incrementar el rendimiento de cultivos productivos de chícharo (*Lathyrus sativus* L.) en suelos con retención de fósforo.

La información sobre el requerimiento de fósforo de cultivos de chícharo (*Lathyrus sativus* L.) es muy escasa, particularmente en suelos que retienen este elemento. En consecuencia, se evaluó el efecto de la fertilización fosfatada sobre el rendimiento y peso del grano de chícharo (variedad Luanco-INIA) en suelos con baja disponibilidad de P y alta capacidad de retención de P, en campos de pequeños agricultores de la Región de La Araucanía, sur de Chile (37°30' -39°30' S). Los ensayos se realizaron durante 2000-2001, 2001-2002 y 2002-2003 en seis sitios; tres en el área de Lumaco y tres en el área de Selva Oscura. Se evaluaron seis dosis de P (0; 21,8; 43,6; 65,4; 87,2 y 109,0 kg ha^{-1}) en un diseño de bloques completos al azar con cuatro repeticiones. Se sembró a razón de 47 semillas m^{-2} . El rendimiento de grano de todos los ensayos promedió 2456 kg ha^{-1} . La fertilización fosfatada incrementó el rendimiento de grano del chícharo durante las temporadas agrícolas 2000-2001 y 2001-2002, en ambas áreas. No hubo efecto significativo en 2002-2003, temporada de cultivo con una caída pluviométrica inusualmente elevada en primavera-verano, lo cual podría haber aumentado la tasa de mineralización de P desde la fracción orgánica del suelo y en consecuencia la disponibilidad de P. Asimismo,

es posible que el crecimiento del sistema radical haya sido favorecido y con ello la exploración de un mayor volumen de suelo. De acuerdo a este estudio, cultivos de chícharo de esta variedad, en suelos con menos de 10 mg kg^{-1} de P-Olsen disponible, deberían responder a la fertilización fosfatada.

Palabras clave: almorta, *Lathyrus*, fósforo, cultivos desatendidos, leguminosas de grano de estación templada fría.

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ORGANIC MATTER REDUCES COPPER TOXICITY FOR THE EARTHWORM *Eisenia fetida* IN SOILS FROM MINING AREAS IN CENTRAL CHILE

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ABSTRACT

The Aconcagua River basin (Central Chile) is one of the most important agricultural areas in the country. However, several copper (Cu) mining operations are located in the basin. The objective of the study was to determine Cu toxicity for the earthworm *Eisenia fetida* (Savigny 1826) in the agricultural soils of the basin. We determined the production of cocoons and juveniles of earthworms in the studied soils. The soils differed in the concentrations of organic matter (OM, range 2-6%), pH (range 7.3-8.3), texture (from loamy sand to clay loam), and total Cu concentrations (range 230-960 mg kg⁻¹). Concentrations of Cu and OM in the soils were the variables that determined the earthworms' biological response. In contrast, pH and texture did not affect this response. Cocoon and juvenile production decreased considerably in soils with elevated Cu concentrations (> 500 mg kg⁻¹), regardless of OM concentrations. Cocoon production decreased in the soils with Cu concentrations below 500 mg kg⁻¹ when OM concentrations were below 3.5%. In contrast, cocoon production did not vary when OM concentrations were above 3.5%. The same effect of OM was observed on juvenile production. In this case, the threshold for OM concentration was 2.5%. It was concluded that it is important to consider OM concentrations in order to predict the biological response of earthworms in these soils.

Key words: *Eisenia fetida*, Aconcagua River, ecological risk assessment, Cu mining, trace elements.

INTRODUCTION

Copper (Cu) mining is the most important economic activity in Chile. However, the environmental problems historically associated with copper mining are widely known, particularly in relation to the contamination of agricultural soils by trace elements such as Cu (González *et al.*, 2008; De Gregori *et al.*, 2003). Although Cu is an essential element for all organisms, it becomes toxic at high concentrations (Sauvé *et al.*, 1998).

The Aconcagua River basin in Central Chile is one of the most important agricultural areas in the country. On the other hand, several copper mining industries are located in the agricultural areas of the basin. There is little information available about the toxicity of copper for organisms and crops in agricultural soils of Chile.

Knowing the total concentration of a trace element in a soil is not sufficient to predict the potential ecological risk that it represents (Sauvé *et al.*, 1998). Ecological risk is more related to the bioavailability of the element that, in turn, is related to the chemical form in which it is found in the soil. The National Research Council (NRC, 2003) defines bioavailability as the fraction of the total element that is available to the receptor organism.

Chile currently does not have any legislation on the maximum acceptable concentrations of toxic elements in soils. In the opinion of the authors based on what is outlined above, any future legislation should distinguish between soils where trace elements are present but do not represent a risk from those that, at similar concentration of trace elements, do represent significant ecological risks.

An approximation that can be used to solve this problem is carrying out toxicity bioassays with soil macroorganisms. Standardized bioassays that determine the acute and chronic toxicity with earthworm *E. fetida* are particularly suitable (OECD, 2000). *E. fetida* is considered representative of soil macrofauna and of earthworms in particular (OECD, 2000). The objective of the present study was to determine the toxicity of trace elements for earthworms in agricultural soils from mining areas in Central Chile.

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MATERIALS AND METHODS

Site selection and soil sampling

The selection of sampling sites was based on the results of a previous study on the distribution of copper in agricultural soils in the Aconcagua River basin (R. Aguilar, unpublished results). This study revealed that the Catemu Creek sub-basin has the largest surface area of soils with high concentrations of Cu.

With the objective of obtaining samples with a wide range of total Cu concentration, 13 localities of the Catemu Creek sub-basin were sampled (Table 1). In each locality, 10 kg of soil were obtained, from a soil depth of 0 to 20 cm, following the removal of the existing vegetation. The soils were then taken to the laboratory of the Faculty of Agriculture, Pontifical Catholic University of Valparaiso, where they were dried at 60 °C for 2 days. Then, the soils were disaggregated in a porcelain mortar and sieved to 2 mm.

Physical-chemical analysis of the soils

The total concentrations of Cu, lead (Pb) and zinc (Zn) were determined by atomic absorption spectrophotometry (GBC, model 902, Dandenong, Victoria, Australia) following acid digestion of the soils with a mixture of fluorhydric and perchloric acids (Maxwell, 1968). The determination of total arsenic (As) in the soils was carried out using neutron activation analysis. To ensure the quality of the results, reference samples were analyzed. For all the cases, the percentage of difference between the obtained values versus the certified values did not exceed 10%.

The soil texture was determined using the simplified hydrometer method according to Sheldrick and Wang (1993). The concentration of organic matter (OM) was determined according to Sadzawka *et al.* (2006). Electrical conductivity (EC) and pH were determined in saturated paste extracts (Sadzawka *et al.*, 2006). Soluble Cu concentration and activity of free Cu²⁺ ion (pCu²⁺) were determined in the same extract by atomic absorption spectrophotometry and by ion selective electrode (Rachou *et al.*, 2007a), respectively.

Bioassays of toxicity

To determine chronic toxicity, bioassays with the earthworm *Eisenia fetida* were carried out using the protocols of the OECD (2000). Specifically, 500 g of soil was adjusted to a humidity of 40% w/w and placed in experimental glass containers of 750 mL. Ten adult earthworms (with visible clitellum) were incubated in each container. Earthworms were previously washed with distilled water, blotted dry and weighed. Five grams of cow manure were moistened with 5 mL of distilled water and added to each container. Eight replicates were made with each soil. The design was randomized. After 4 weeks of exposure, the weight, adult survival and number of cocoons were determined. Then, the cocoons were incubated in the same soil for additional 4 weeks and the number of juveniles was determined. Room temperature was maintained within the range of 22 to 24 °C, with illumination of 200 lux and a photoperiod of 12 h of light and 12 h of darkness. Moisture was maintained by the application of 40 mL of distilled water once a week. Soil 41 was considered as a control because of the lowest concentrations of Cu among the studied soils (Table 1).

Table 1. Physico-chemical characteristics of the studied soils.

Soil	Texture	OM	Cu	Pb	Zn	As	CE	pH	pCu ²⁺	Cu
		%	mg kg ⁻¹				dS m ⁻¹			mg L ⁻¹
18	Sandy loam	5.2	959	542	923	44	2.3	7.3	7.3	2.70
27	Sandy loam	3.5	847	81	209	19	2.4	7.9	7.9	0.74
28	Loam	2.2	382	81	244	17	1.3	8.0	8.0	0.28
29	Loam	4.0	431	30	156	21	0.8	8.3	8.3	0.67
30	Clay loam	5.2	426	46	135	30	2.2	7.6	7.6	0.29
31	Loam	5.3	354	36	117	24	2.9	8.1	8.1	0.56
37	Loamy sand	2.0	354	36	97	30	2.6	7.9	7.9	0.38
38	Loam	3.5	434	40	117	28	2.5	7.9	7.9	0.22
41	Loam	6.0	226	36	148	29	4.8	7.8	7.8	0.60
49	Loam	5.6	707	47	134	37	4.1	8.0	8.0	0.83
50	Clay loam	4.0	650	63	146	45	1.5	8.2	8.2	0.46
52	Loam	2.8	597	47	133	36	2.1	8.2	8.2	0.48

OM: organic matter. CE: electrical conductivity. pCu²⁺: -log (activity of the free Cu²⁺ ion).

Statistical analysis

Using the Dunnett test, comparisons between the responses of the earthworms in the studied soils and the control soil have been made. Simple and multiple regressions were made between the responses of the earthworms and the physico-chemical characteristics of the soils. Also, simple regressions were made between the concentrations of Pb, Zn or As and those of Cu. We used the Minitab 3.1 and Excel 2003 for statistical analysis.

RESULTS AND DISCUSSION

Soil characterization

The studied soils were different in OM concentrations (range of 2 - 6%), EC (range of 0.8 - 4.8 dS m⁻¹), pH (range of 7.3 - 8.3) and texture (from loamy sand to clay loam). The soils presented a wide range of total Cu concentration (from 230 to 960 mg kg⁻¹) (Table 1). These high Cu concentrations are mainly due to mining activities, while application of copper-based products in agriculture represents a minor source (R. Aguilar, unpublished results). Simple regressions revealed that the relations between the concentrations of Pb, Zn and As (Table 1) versus those of Cu were not significant ($P > 0.05$).

Validity of the control soil

The OECD (2000) established the following criteria for validity of the control soil: (1) each repetition (10 adult earthworms) should produce at least 30 juveniles at the end of the bioassay, (2) the variation coefficient in the reproduction parameters must be less than 30%, and (3)

adult mortality must be less than 10%. Additional to this, Spurgeon *et al.* (2003) propose that weight loss should be less than 15%. These four criteria were satisfied in the control soil (Table 2).

Survival and weight loss

In all the soils used, the earthworm survival was higher than 98% and weight loss did not exceed 20% (Table 2). The data of survival presented a very narrow range (98-100%) and, thus, it was not possible to carry out regressions with the physico-chemical characteristics of the soils. On the other hand, the physico-chemical characteristics of the soils did not explain weight variation. As shown below, the survival and weight loss are variables that less sensitive to Cu toxicity, in comparison to the reproduction variables.

Identification of the variables that affected reproduction

Bioassays with earthworms have been widely used to determine the toxicity of trace elements in soils (Spurgeon *et al.*, 2003). However, the technique has been criticized for not adequately representing real environmental conditions and, consequently, not being relevant from an environmental point of view (Davies *et al.*, 2003). The criticism is based on the fact that the OECD (2000) proposes the use of artificial soils (composed of peat, clay, and sand) enriched with solutions of metals at increasing concentrations. In fact, it has been observed that the toxicity of trace elements for earthworms is considerably higher in artificially-contaminated soil media than in field-collected soils. This is explained by a greater bioavailability of trace

Table 2. Results of chronic toxicity bioassays (OECD, 2000). Soil 41 was considered as a control (C).

Soil	Survival	Weight loss	Number of cocoons	Number of juveniles	Cu in earthworms
	%				
41 (C)	100 ± 0	(-)14 ± 3.8	28 ± 3.2	54 ± 4.4	23 ± 0.8
18	100 ± 0	(-) 4 ± 8.9	11 ± 6.5*	19 ± 17*	61 ± 31*
27	99 ± 3.5	(-) 8 ± 5.4	6 ± 4.0*	14 ± 9.0*	64 ± 4.4*
28	100 ± 0	(-)10 ± 9.7	9 ± 5.6*	8 ± 9.8*	37 ± 0.4
29	100 ± 0	(-)15 ± 3.5	26 ± 5.2	50 ± 12	37 ± 3.8
30	98 ± 4.6	(-) 8 ± 5.8	18 ± 3.0*	36 ± 4.4*	39 ± 1.9
31	100 ± 0	(+) 1 ± 3.9*	20 ± 4.1*	41 ± 5.7	51 ± 4.5*
37	100 ± 0	(-) 8 ± 4.5	15 ± 6.3*	23 ± 7.9*	41 ± 2.9
38	99 ± 3.5	(+) 2 ± 6.8*	15 ± 5.5*	41 ± 17	59 ± 3.5*
49	100 ± 0	(-)19 ± 8.0	9 ± 5.1*	24 ± 12*	46 ± 0.6
50	98 ± 4.6	(-) 6 ± 6.8	4 ± 3.1*	14 ± 7.8*	46 ± 0.6
52	100 ± 0	(+) 2 ± 12*	5 ± 4.3*	13 ± 6.8*	37 ± 27

* Significantly different from the control according to the Dunnett test ($P < 0.05$).

±: Standard deviation. (-) = Weight loss. (+) = Weight increase.

elements in artificially-contaminated soils in comparison to those collected in the field (Spurgeon and Hopkin, 1995). As a result, recent studies highlight the importance of using field-collected soils to evaluate ecological risk of trace elements present in the soil (Nahmani *et al.*, 2007a; 2007b).

On the other hand, the use of field-collected soils presents several difficulties. First, in areas near copper mining activities, soils have high concentrations of several trace elements (Cu, Pb, Zn, Cd and As, among others; De Gregori *et al.*, 2003; Ginocchio *et al.*, 2004). In this case, it could be difficult to distinguish between the effects of different trace elements on the response of the earthworms. Second, agricultural soils can contain other types of chemical compounds, such as pesticides and/or fungicides, which can affect the response of the earthworms (Slimak, 1997). Finally, the intrinsic physico-chemical characteristics of the soil, such as pH, texture and OM content, among others, also affect the degree of toxicity of the trace elements present in the soil (Kennette *et al.*, 2002, Nahmani *et al.*, 2007a).

In the present study, the variables that affected earthworm reproduction were identified using simple and multiple regressions between earthworm responses and the physico-chemical characteristics of the soils. The regressions ruled out any evident effects of Pb, Zn, and As on the response of earthworms. The simple and multiple regressions between earthworm reproduction and the physico-chemical characteristics of the soils indicated that pH, texture, EC, and soluble Cu concentration did not affect the response of the earthworms. On the other hand, a significant regression was observed between pCu^{2+} and earthworm reproduction (Table 3). The effect of free Cu^{2+} ion will be discussed in detail below.

The best prediction of the earthworm response was obtained by considering total Cu concentrations together with OM (Table 3). The regression coefficients increased upon considering both variables together, in comparison to total Cu alone. This effect of OM on Cu toxicity will be discussed in detail below.

Thus, the effects observed on earthworm reproduction are mainly due to Cu and OM, explaining about 70% of the variance (Table 3). Nevertheless, the studied soils could contain other undetermined compounds (for example, pesticides and/or fungicides) that have affected earthworm reproduction.

Effect of OM on the toxicity of copper

The earthworm *E. fetida* lives in environments rich in OM (OECD, 2000). Despite this, the regressions between OM concentrations and earthworm reproduction were not significant. This suggests that OM does not have a direct effect on the reproduction parameters. This concurs with Spurgeon and Hopkin (1999) who indicated that *E. fetida* was not able to obtain sufficient nutrients from mineral soils (with OM < 20%), requiring the addition of food to the soils used in the bioassays.

The multiple regressions show that OM promotes cocoon and juvenile production, while total Cu decreases earthworm reproduction. Consequently, it is necessary to consider OM content to predict the biological responses of earthworms in soils contaminated with trace elements.

The soils with more than 50% of inhibition in the production of cocoons or juveniles are considered as toxic for earthworms (Hund-Rinke and Wiechering, 2001; Hund-Rinke *et al.*, 2005). The soils with total Cu concentrations higher than 500 mg kg^{-1} were toxic, independent of the OM concentrations (Figure 1). In contrast, in the soils with total Cu concentrations below 500 mg kg^{-1} , OM concentrations determined Cu toxicity. In the case of cocoon production, the soils with total Cu concentrations below 500 mg kg^{-1} were toxic when OM concentrations were lower than 3.5%. The opposite was observed in the case of OM concentrations above 3.5% (Figure 1). In the case of juvenile production in soils with total Cu concentrations below 500 mg kg^{-1} , the critical threshold for OM concentration was about 2.5% (Figure 2).

The combined effect of total Cu and OM on the response of earthworms is due to the control that these two variables exert on the activity of the Cu^{2+} ion that

Table 3. Regressions between the number of cocoons/juveniles and soil copper concentrations.

Production of cocoons	R ²	P	Production of juveniles	R ²	P
PC = 26 + 0.02 CuT	0.44	0.02	PJ = 50 - 0.04 CuT	0.34	0.05
PC = 15 - 1.6 CuS	0.21	0.12	PJ = 30 - 3 CuS	0.23	0.10
PC = 16 - 0.03 CuT + 2.8 OM (0.005) (0.03)	0.68	0.01	PJ = 24 - 0.05 CuT + 6.8 OM (0.01) (0.01)	0.68	0.01
PC = -69 + 8.5 pCu^{2+}	0.43	0.02	PJ = -118 + 15 pCu^{2+}	0.35	0.04

PC: Production of cocoons. PJ: production of juveniles. OM: organic matter (%). CuT: total copper (mg kg^{-1}). CuS: soluble copper (mg L^{-1}). pCu^{2+} : -log (activity of the free Cu^{2+} ion). R²: regression coefficient. P: probability. Number in parenthesis indicates the P value of the variable in multiple regressions.

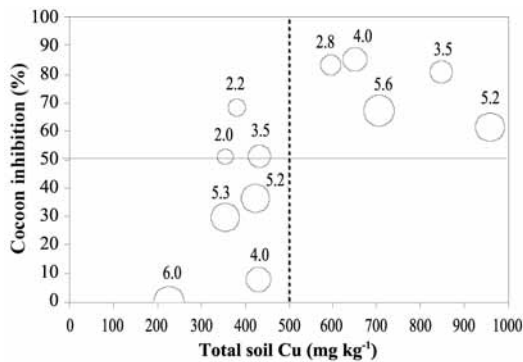


Figure 1. Effect of total copper and organic matter on cocoon production. The size of the circles and the values represent the percentage of organic matter in each soil.

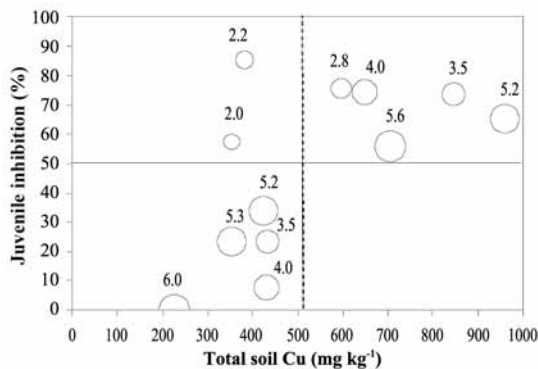


Figure 2. Effect of total copper and organic matter on juvenile production. The size of the circles and the values represent the percentage of organic matter in each soil.

is considered as the bioavailable form of Cu in the soil (Sauvé *et al.*, 1998; Thakali *et al.*, 2006). McBride *et al.* (1997) and Sauvé *et al.* (1997) proposed an empirical equation that describes the effect of the physico-chemical properties of the soil on the activity of the Cu^{2+} ion: $\text{pCu}^{2+} = a + b \text{pH} + c \log \text{CuT} + d \log \text{OM}$. In agreement with this postulate, in the present study, the activity of the Cu^{2+} ion in the saturated paste extracts was controlled by the concentrations of OM and total Cu (Table 4). In turn, pH did not affect pCu^{2+} , probably because of the narrow pH range in the studied soils (pH of 7.3 to 8.3).

Also, the effect of OM in reducing Cu toxicity for earthworms is probably due to a change in the mobilization of copper from the solid phase to the soil solution. According to Rachou *et al.* (2007b), the kinetics of the mobilization of the elements from the solid phase to the soil solution decreases with increasing concentration of OM. Thus, the decrease in the flow of Cu from the solid phase to the soil solution can, in turn, reduce its toxicity for the earthworms.

Effect of different forms of copper on reproduction

The bioavailability of a trace element can be estimated through a chemical analysis that extracts a fraction of the element. Diverse extractants were proposed to simulate the bioavailability of trace elements for plants and soil organisms. It is often considered that the bioavailable fraction of a trace element corresponds to its soluble form (Posthuma *et al.*, 1997; Kabata-Pendias, 2004). Nevertheless, the regressions between earthworm reproduction and Cu concentrations in the saturated paste extract, corresponding to the soluble form of Cu (Table 3), indicated that this form of the element did not affect the earthworms' reproduction.

On the other hand, the Cu^{2+} ion is the bioavailable form of Cu both in soil and water (Thakali *et al.*, 2006). The activity of the Cu^{2+} ion is often considered to be the best variable to predict Cu toxicity for plants, organisms and microbial processes in the soil (Sauvé *et al.*, 1998). In accordance with these postulates, in the present study, the regressions between pCu^{2+} in saturated paste extract and earthworms' reproduction were significant (Table 3).

Bioaccumulation of copper

Earthworms can actively excrete Cu assimilated in their tissues (Spurgeon and Hopkin, 1999). This implies extra energy costs, generating a reduction in energy available for growth and development. This, in turn, affects sexual maturity and production of cocoons and juveniles (Spurgeon and Hopkin, 1996). Likewise, the efficiency of excretion probably decreases with increasing concentrations of Cu, resulting in increased bioaccumulation (i.e., the concentration of assimilated Cu in earthworm tissue) in the soils with higher Cu concentrations (Svendsen and Weeks, 1997; Scott-Fordsmand *et al.*, 2000).

In the present study, the concentrations of total Cu in the soil explained 53% of the variance ($P = 0.001$) in the bioaccumulation of Cu. In contrast, other soil properties like pH, OM and other forms of Cu (soluble or free) did

Table 4. Effect of total copper, organic matter and pH on pCu^{2+} .

	R^2	P
$\text{pCu}^{2+} = 15 - 2 \log \text{CuT}$	0.34	0.04
$\text{pCu}^{2+} = 14 - 2 \log \text{CuT} + 1.9 \log \text{OM}$ (0.018) (0.042)	0.59	0.01
$\text{pCu}^{2+} = 7 - 2 \log \text{CuT} + 2 \log \text{OM} + 0.8 \text{pH}$ (0.020) (0.014) (0.093)	0.72	0.01

CuT: total copper (mg kg⁻¹). OM: organic matter (%). pCu^{2+} : -log (activity of the free Cu^{2+} ion).

R^2 : regression coefficient. P: probability. Numbers in parenthesis indicate the P value of the variable in multiple regressions.

not affect its bioaccumulation. The concentrations of Cu in the tissues were in the range of 23 to 64 mg kg⁻¹. The effect of these concentrations on the earthworm reproduction is discussed in detail below.

The normal range of Cu concentration in earthworm tissues can be determined through the use of biomarkers of the stress induced by this element. For example, Svendsen and Weeks (1997) used the stability of the lysosomal membrane as a biomarker of sub-cellular stress in the earthworms *E. andrei* exposed to increasing concentrations of Cu. The degree of damage induced by Cu on the lysosomal membrane depended on its bioaccumulation. No damage to tissue was detected at concentrations of 8 to 25 mg kg⁻¹. Concentrations of 25 to 55 mg kg⁻¹ produced medium damage, while concentrations higher than 55 mg kg⁻¹ provoked severe damage. Likewise, Scott-Fordsmann *et al.* (2000) reported the critical threshold of 50 mg kg⁻¹ of bioaccumulation of Cu by *E. fetida*, using the same biomarker of damage to the lysosomal membrane. Similarly, the species *Lumbricus rubellus* did not show a decrease in the production of cocoons with a Cu bioaccumulation below the critical threshold of 40 mg kg⁻¹ (Ma, 2005).

In the present study, the bioaccumulation of 23 mg Cu kg⁻¹ in the earthworms present in the control soil can be considered as normal, in accordance with Svendsen and Weeks (1997). A higher bioaccumulation of Cu (in the range of 37 to 64 mg kg⁻¹) caused a reduction in cocoon production. Svendsen and Weeks (1997) proposed that earthworms can present individual differences in the efficiency of excreting assimilated Cu. Consequently, bioaccumulation of Cu is not a good biomarker of stress induced by this element, as is reflected in the low regression coefficient ($R^2 = 0.37$, $P < 0.05$) between the parameters of reproduction and bioaccumulation.

CONCLUSIONS

The majority of the studied soils in the Aconcagua River basin in Central Chile presented toxic effects on earthworms, inhibiting the production of cocoons and juveniles.

The regression analysis ruled out any evident effects of Pb, Zn, and As on the response of earthworms.

The observed effects on earthworm reproduction are mainly due to Cu and OM, explaining 70% of the variance.

Reproduction of earthworms is not determined solely by Cu, but also by OM. It is necessary to know OM concentrations to correctly predict the response of macrofauna in soils contaminated by Cu.

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RESUMEN

Materia orgánica reduce la toxicidad del cobre para la lombriz *Eisenia fetida* en suelos de áreas mineras en Chile Central. La cuenca del Río Aconcagua (Chile Central) es una de las más importantes áreas agrícolas en el país. Por otro lado, varias industrias de la minería de cobre (Cu) se encuentran ubicadas en esta cuenca. El objetivo del estudio fue determinar la toxicidad de Cu para la lombriz *Eisenia fetida* (Savigny 1826) en los suelos agrícolas de la cuenca. Se determinó la producción de capullos y juveniles de la lombriz en suelos estudiados. Los suelos se diferenciaron por las concentraciones de materia orgánica (MO, rango 2-6%), pH (rango 7,3-8,3), textura (entre arenoso franca y franco arcillosa) y concentraciones totales de Cu (rango 230-960 mg kg⁻¹). Las concentraciones de Cu y MO en los suelos fueron las variables que determinaron la respuesta biológica de las lombrices. En contraste, pH y textura no afectaron a esta respuesta. La producción de capullos y juveniles disminuyó considerablemente en suelos con altas concentraciones de Cu (> 500 mg kg⁻¹), independientemente de las concentraciones de MO. La producción de capullos disminuyó en suelos con concentraciones de Cu inferiores a 500 mg kg⁻¹ cuando las concentraciones de MO fueron inferiores a 3,5%. Por el contrario, la producción de capullos no varió cuando la concentración de MO fue superior a 3,5%. El mismo efecto de MO fue reconocido sobre la producción de juveniles. En este caso, el umbral crítico de la concentración de MO fue de 2,5%. Se concluye la importancia de considerar las concentraciones de MO para predecir las respuestas biológicas de lombrices en estos suelos.

Palabras clave: *Eisenia fetida*, Río Aconcagua, evaluación del riesgo ecológico, minería de Cu, elementos traza.

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GENE EXPRESSION ANALYSIS: A WAY TO STUDY TOLERANCE TO ABIOTIC STRESSES IN CROPS SPECIES

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ABSTRACT

Regions traditionally destined to agriculture report an ever increasing exposure to cold and drought conditions. This is especially important in countries like Chile where crop management options are limited. The development of new cultivars with better yields under adverse conditions is fundamental if the ever increasing demand for food is to be matched; however, improving tolerance to abiotic stresses has proved to be a complex task. In this regard, development in plant physiology and genomics in the last 20 years has led to a deeper understanding of how plants respond to stress and mechanisms responsible for different ranges of tolerance observed in nature. This review discusses the techniques currently most in use in gene expression analysis, together with some important experimental design variables, such as the developmental stage of the plant, stress intensity and duration, and how different stresses may interact when performing assays. On the other hand, it is fundamental to properly select gene expression techniques according to the available information on the genome, the crop and the final objective of the research. All these points must be considered to ease transition from genomics to practical applications to crop species in order to increase their tolerance to stress. In this regard, the rapid development of new techniques in gene expression analysis with lower costs will determine a new revolution in crop research in coming decades. Therefore, Chile needs to be prepared in this area to continue its development as a major food producer worldwide.

Key words: cold, drought, crops species, gene expression.

INTRODUCTION

Episodes of low or high temperature and drought are among the environmental conditions that most plants experience on a daily basis. In crops, this variation from ideal growth conditions often results in lower yields and a high economic impact for producers and consumers.

Understanding the mechanisms involved in the response of plants to adverse environmental conditions is, without a doubt, the first step in the generation of crops with higher tolerance to stress. Research at the level of genes (genomics), proteins (proteomics), metabolites (metabolomics), individuals (physiology, systemic- biology) and communities (ecology) has been fundamental in the current understanding of the response of plants to stress. In particular, a huge development in the field of genomics in the last 20 years has led to a deeper understanding in areas such as gene expression, organization and its relationship to stress tolerance. Functional genomics studies the function of genes of an organism and focuses

on dynamic processes such as transcription, translation, interaction of genes and how they are related to different phenotypes. Connecting gene function and traits relevant to agriculture, such as yield, plant structure and tolerance to adverse environmental conditions has become of utmost interest considering global warming, urban development and an ever increasing population demand for food.

Genome analysis has been mostly limited to model plants that fulfil some specific requirements such as: (1) small genome size, (2) short generation time, (3) small size to enable growth in limited space, and (4) availability of gene manipulation technologies (Tabata, 2002). In particular, two of the most important model species are *Arabidopsis thaliana* and rice (*Oryza sativa* L.) for dicotyledonous and monocotyledoneous plant species, respectively. Besides its importance as a crop, rice has a high degree of synteny with genomes of other cereals plants, such as maize, wheat, barley and other grasses because their genomes share a considerable similarity in their organization, as well as sequence similarity (Gale and Devos, 1998; Bowers *et al.*, 2005; Paterson *et al.*, 2005). Great advances in the comparison of genomes and transcriptomes of different organisms have contributed to the development of comparative genomics as one of the most promising fields in the area (Gale and Devos, 1998;

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Caicedo and Purugganan, 2005). In this way, finding variations in the genome or the transcriptome from the current model species related to interesting agronomic traits is of the highest importance for crop biotechnology (van de Mortel and Aarts, 2006).

The objective of this review is to summarize current techniques used in gene expression analysis in plants and their relevance to abiotic stress research. Special emphasis is given to issues to be considered when comparing performance of crops in controlled conditions and in the field.

Techniques used for evaluating gene expression in functional genomics studies

A fundamental step in any functional genomics study is the analysis of gene expression. One of the greatest strengths of genomics compared to other disciplines is the prospect of analyzing the expression of thousands of genes simultaneously, resulting in a more comprehensive picture of changes occurring in the transcriptome across different conditions (Green *et al.*, 2001).

The technology available for the analysis of gene expression can be divided into two categories: closed and open systems (Table 1). Closed systems are characterized by a finite number of genes that can be assessed by virtue of their inclusion by selection. Therefore, the coverage of genes will be related to the completeness of the knowledge of the genome being studied, limiting this kind of analysis

to the most well characterized species or systems (Green *et al.*, 2001). Typically, closed systems such as microarrays (Table 2) and real-time polymerase chain reaction (PCR) have been extensively used in gene expression analysis in plants (Ma *et al.*, 2005; Rensink *et al.*, 2005; Oono *et al.*, 2006; Xu and Shi, 2006; Mantri *et al.*, 2007; Monroy *et al.*, 2007; Fernandez *et al.*, 2008; Remans *et al.*, 2008). On the other hand, with open systems there is no need for previous knowledge of the genome or transcriptome of the organism. cDNA-AFLP (cDNA-Amplified fragment length polymorphism), MPSS (massively parallel signature sequencing), and specially SAGE (serial analysis of gene expression) have been successfully used to quantify transcript abundance and generate expression data across different tissue types or developmental stages in higher plants (Fizames *et al.*, 2004; Meyers *et al.*, 2004; Calsa and Figueira, 2007; Chen *et al.*, 2007; Leymarie *et al.*, 2007; McIntosh *et al.*, 2007; Song *et al.*, 2007; Ritter *et al.*, 2008). Worth mentioning are 454-sequencing technology and digital gene expression (DGE) that have recently been used to study the transcriptome of different organisms and promise to become an efficient and cost-effective alternative with high potential in crop research (Mikkilineni *et al.*, 2004; Margulies *et al.*, 2005; Velculescu and Kinzler, 2007; Weber *et al.*, 2007; Torres *et al.*, 2008). Open and closed systems should not be considered as competitors, but rather as complementary technologies to

Table 1. Most commonly used techniques for gene expression analysis in plants.

Open systems	<p>cDNA-AFLP. RNA is converted into double stranded cDNA and then digested with two restriction enzymes: a frequent-cutter and a rare-cutter. Synthetic adapters are ligated to the cDNA ends and primers complementary to the adapter sequences (plus small extensions of 1, 2, or 3 nucleotides) are used to amplify fragments with asymmetric ends. These fragments are displayed on sequencing gels and compared. Specific fragments can be eluted from gels and sequenced to identify genes with differential expression (Bachem <i>et al.</i>, 1996).</p>
	<p>SAGE. RNA is converted to double-stranded cDNA with a biotin attached to the oligo(dT) first strand synthesis primer and cleaved with a restriction enzyme, leaving 3'-most fragments immobilized onto streptavidin beads. After ligation with linkers onto the non-biotinylated end cDNAs are released, ligated together and amplified by PCR. Primer regions are removed from PCR products and the resulting fragments are ligated together into concatemers, cloned and sequenced. Finally, a software package identifies and counts the relative frequency of the sequences in the samples (Velculescu <i>et al.</i>, 1995; Yamamoto <i>et al.</i>, 2001).</p>
	<p>MPSS. Individual 3' restriction fragments from a cDNA library are coupled to one of a million beads, amplified, arrayed and sequenced simultaneously for 20 residues to provide a million signature sequences. Transcripts can then be identified and the corresponding transcriptome quantitatively characterized (Brenner <i>et al.</i>, 2000).</p>
Closed systems	<p>Microarrays. Marked samples are tested against sequences from thousands of different genes fixed on small solid supports (usually glass microscope slides). Depending on their sequence, the samples will hybridize with different spots in the array, which is analyzed by specialized image software (Schena <i>et al.</i>, 1995).</p>
	<p>Real-time PCR. A variant from conventional PCR based on the detection and quantification of the fluorescence emitted by PCR products accumulated through the amplification process (Higuchi <i>et al.</i>, 1993).</p>

be used depending on the subject to be analyzed and the objectives of the research.

The amount of material available is an important variable to be considered in the selection of a technology for gene expression analysis. In particular, the sensitivity and coverage of the method will be determinant considering that 90-95% of all mRNA species are present at five or fewer copies per cell (Green *et al.*, 2001). When insufficient quantities of RNA are obtained, cDNA generated from RNA can be amplified exponentially by PCR, or linearly with T7 RNA polymerase to avoid differential amplification (Brady *et al.*, 2006).

Another variable to keep in mind is the selection of the sample to be analyzed. In this regard, the latest advances in microdissection techniques allow extraction of RNA from specific tissues and individual cells, opening the possibility of a highly detailed analysis and a considerable reduction of noise generated by the natural heterogeneity of plant organs (Brandt, 2005; Lee *et al.*, 2005; Ohtsu *et al.*, 2007).

Abiotic stress variables to be considered in functional genomics studies

The aim of most functional genomics studies concerned with abiotic stress is to relate gene function to traits of plant performance under adverse environmental conditions.

A recurring question is how representative are growth chamber studies compared to field studies. In this respect there is a lack of studies that comprehensively evaluate correlations between growth chambers and the field in terms of plant performance. The issues in what follows should be considered when designing an experiment in controlled conditions with possible applications in the field:

Combination of stresses. Most studies so far have been focused on the response to just one kind of stress. This strategy has led to key discoveries that otherwise would not have been possible and that have helped us to understand in greater depth the way plants respond to stress. However, it should be noted that plants in the field are usually exposed to more than one stress simultaneously. This combination of stresses is fundamental to understand differences between the performance of crops in controlled growth chambers and in the field (Knight and Knight, 2001; Mittler, 2006).

Length of the treatment. Despite the fact that valuable data can be obtained from short term experiments, it is longer term plant performance with respect to biomass, yield data and the degree of recovery from stress that has the most value in agriculture (Vinocur and Altman, 2005).

Table 2. Gene expression analysis through microarrays in some crops.

Plant	Stress	Reference
<i>Capsicum annuum</i>	Cold stress	Hwang <i>et al.</i> , 2005
<i>Hordeum vulgare</i>	Drought and salinity	Ozturk <i>et al.</i> , 2002
<i>Oryza sativa</i>	Cold, drought, salinity and ABA ¹ treatment	Rabbani <i>et al.</i> , 2003
	Chilling stress	Yamaguchi <i>et al.</i> , 2004
	Drought stress	Hazen <i>et al.</i> , 2005
	Drought stress	Lan <i>et al.</i> , 2005
	Drought stress	Wang <i>et al.</i> , 2007
	Drought and high salinity	Zhou <i>et al.</i> , 2007
<i>Solanum tuberosum</i>	Cold, heat and salt stress	Rensink <i>et al.</i> , 2005
<i>Sorghum bicolor</i>	Dehydration, salt and ABA treatment	Buchanan <i>et al.</i> , 2005
<i>Triticum aestivum</i>	Low-temperature stress	Gulick <i>et al.</i> , 2005
	Drought	Mohammadi <i>et al.</i> , 2007
<i>Helianthus annuus</i>	Chilling stress	Fernandez <i>et al.</i> , 2008
<i>Zea mays</i>	Salinity	Wang <i>et al.</i> , 2003
<i>Manihot esculenta</i>	Heat and drought	Sakurai <i>et al.</i> , 2007

¹ABA: amino butyric acid.

Intensity of the treatment. Plants respond in different manners to variable degrees of stress, as different protection/repair mechanisms will be engaged accordingly. For example, the response of a plant to chilling stress will be different from the response to freezing stress, considering that the latter can lead to ice formation. The intensity of the treatment will also be of the utmost importance for screening purposes, especially when a ranking of tolerance to stress is to be established.

Stage of crop development. Clearly, tolerance to stress is different throughout the lifecycle of any plant and the consequences of exposure to stress may also vary. As an example, rice is especially susceptible to low temperature during the germination and reproductive stages (Board *et al.*, 1980; Jacobs and Pearson, 1994) In the first, a possible consequence is the failure to germinate. Exposure to cold during the reproductive stage will induce sterility rather than have an effect on plant survival. In this regard, it is important to have plants in similar stages of development when screening for tolerance to stress, especially with cultivars that complete their lifecycle at different times.

Designing experiments in functional crop genomics must consider all these recommendations in order to successfully extrapolate results to the field. As well, it is important to keep in mind genotype x environment interactions (G x E) when evaluating the performance of any genotype in the field by including the range of adaptation of new varieties to different environments and the consistency of their performance over time. In order to effectively recognize G x E in any breeding program, there must be a comprehensive characterization of the genotypes and environments being assayed, and these considerations are valid for genomics as well.

Functional genomics and stress response in crops

Abiotic stresses are estimated to reduce yield to less than half compared to the potential under ideal growing conditions (Boyer, 1982). Unlike plant resistance to biotic stresses, which is mostly monogenic, tolerance to abiotic stresses are generally multigenic, quantitative and complex traits controlled by quantitative trait loci (QTL). This has clear consequences for the development of plants that are more tolerant to abiotic stresses by genetic engineering (Vinocur and Altman, 2005). A further complication is that some genes may exert control over different traits, resulting in unwanted changes in agronomic plant traits.

Cold and drought tolerance in crops constitute highly desired traits in Chile given the economic consequences of the current climatic trend of very low temperatures in winters and severe drought in summers. Cold stress in plants causes a reduction in enzyme activities, reaction rates, energy imbalance and is accompanied by changes

in the transcriptome, proteome and metabolome (Guy *et al.*, 2008). On the other hand, when plants are exposed to drought, there is a characteristic response of a partial-to-total stomatal closure, resulting in a reduction of CO₂ uptake, transpiration and a major impact for photosynthesis and source-sink relationships (Chaves *et al.*, 2002).

The consequences of any stress will depend on its intensity. As an example, chilling temperatures will be responsible for lower metabolic rates and energy imbalance, while freezing temperatures will additionally cause membrane injury and severe dehydration when ice forms (Graham and Patterson, 1982; Thomashow, 1998; Pearce, 2001).

Drought and temperature stress might occur alone or in combination at any stage in plant development, causing reduced grain weight and yield loss (Sreenivasulu *et al.*, 2007). It is known that exposure to one kind of stress usually involves an increased tolerance to other stresses given that similar effects are shared at the cellular level. As an example, freezing temperatures, low water availability and high salinity can all cause lowering of the cellular osmotic potential and thereby activate osmotic stress responses (Langridge *et al.*, 2006). In this regard, it is not unexpected to find promoters that have sequences for transcription factors involved in drought, salt and cold response, suggesting points of convergence at the molecular level (Knight and Knight, 2001). These results, added to a high overlapping of genes involved in the response to cold, drought and high salinity, suggest an intricate coordination of the response to multiple stresses in plants at molecular level (Kreps *et al.*, 2002; Seki *et al.*, 2002; Seki *et al.*, 2004; Matsui *et al.*, 2008).

At first glance, a shared regulatory network involved in the response to multiple stresses opens possibilities for the development of multiple-stress-tolerant plants. However, it must not be forgotten that the combination of some stresses might require conflicting or antagonistic responses. In this way, the acclimation of plants to this combination would require an appropriate response to each individual stress, as well as compensation and adjustment for some of the antagonistic aspects involved (Mittler, 2006; Rizhsky *et al.*, 2004). As an example, when plants are exposed to drought, their stomata are closed, which is clearly an antagonistic response if the plant is simultaneously exposed to heat, when transpiration is necessary to reduce leaf temperature.

Higher tolerance to abiotic stress could be achieved by increasing protective mechanisms (antioxidants, non-photochemical quenching, etc.) or by increasing the capacity to repair the damage caused by stress. In this matter, the capacity of recovery from stress is usually overlooked, despite its relevance considering that cycles of stress and recovery are common under natural

conditions and may have a major impact in yield (Vinocur and Altman, 2005).

In model species, such as *Arabidopsis*, more than 40-50% of identified stress-responsive gene functions remains to be characterized (Sreenivasulu *et al.*, 2007). In this regard, a successful approach in determining gene function comes from sequence comparison with databases and, more recently, the use of coexpression modules with promising results (Subramanian *et al.*, 2005).

Comparative genomics constitute an increasingly important field in order to understand how similar model species and crops are, and how to transfer knowledge obtained from model species to applications in agriculture (Paterson *et al.*, 2005; van de Mortel and Aarts, 2006). In this matter, the choice of putative candidate genes is facilitated by the conservation of gene sequences, order and distribution among species and the existence of similar functional gene categories in morphologically similar organs (Brady *et al.*, 2006; Pflieger *et al.*, 2001).

Changes in the transcriptome among related species under stress reported by different groups are usually hard to compare since treatments are usually performed with different tissues, exposure times, intensities, and using

different technologies. In this way, a careful experimental design with related plants that present different degrees of tolerance to stress can be extremely informative. A successful example is the comparison of the transcriptome of winter and spring wheat, cultivars with different tolerance to cold, exposed to low temperature. This study reports the correlation of gene expression kinetics with tolerance to low temperature, a subject usually overlooked that emphasizes the importance of sampling in functional genomics studies (Gulick *et al.*, 2005; Monroy *et al.*, 2007).

Gene expression profiling has allowed the identification of hundreds of genes induced when plants are exposed to stress (Kreps *et al.*, 2002; Oono *et al.*, 2006; Jianping and Suleiman, 2007; Mantri *et al.*, 2007). The availability of the complete genome sequence of some model plants, such as *O. sativa* and *A. thaliana*, has allowed the development of whole genome tiling microarrays. This constitutes a new powerful technology that has already made possible the identification of several unannotated transcripts responsive to abiotic stress (Gregory *et al.*, 2008; Matsui *et al.*, 2008). However, finding a gene responsive to stress does not necessarily guarantee its participation in

Table 3. Maize and rice transgenics and stress tolerance.

Plant	Gene	Result	Reference
Maize	<i>NPK1</i> (tobacco MAPKKK)	Drought and freezing tolerance	Shou <i>et al.</i> , 2004
	<i>ZmNF-YB2</i> (maize nuclear factor YB2)	Drought tolerance	Nelson <i>et al.</i> , 2007
	<i>ZmPLC1</i> (phospholipase C 1)	Drought tolerance	Wang <i>et al.</i> , 2008
	<i>TsVP</i> (vacuolar-H ⁺ -pyrophosphatase)	Drought tolerance	Li <i>et al.</i> , 2008
Rice	<i>HVA1</i> (Barley group 3 LEA protein)	Drought and salt tolerance	Xu <i>et al.</i> , 1996
	<i>GPAT</i> (Arabidopsis glycerol-3P-acyltransferase)	Chilling tolerance	Yokoi <i>et al.</i> , 1998
	<i>OsCDPK7</i> (rice calcium-dependent protein kinase)	Cold, drought and salt tolerance	Saijo <i>et al.</i> , 2000
	<i>Dadc</i> (<i>D. stramonium</i> arginine decarboxylase)	Drought tolerance	Capell <i>et al.</i> , 2004
	<i>ABF3</i> (Arabidopsis ABRE-binding factor 3)	Drought tolerance	Oh <i>et al.</i> , 2005
	<i>DREB1A</i> (Arabidopsis DRE-binding protein 1)	Drought and salt tolerance	Oh <i>et al.</i> , 2005
	<i>MnSOD</i> (pea Mn superoxide dismutase)	Drought tolerance	Wang <i>et al.</i> , 2005
	<i>SNAC1</i> (rice stress responsive NAC1)	Drought and salt tolerance	Hu <i>et al.</i> , 2006
	<i>OsDREB1</i> (rice DRE-binding protein 1)	Drought, salt and cold stress tolerance	Ito <i>et al.</i> , 2006
	<i>HvCBF4</i> (barley C-repeat binding factor)	Drought, salt and cold stress tolerance	Oh <i>et al.</i> , 2007
	<i>OsCIPK03</i> (rice calcineurin B-like protein-interacting protein kinase 03)	Cold tolerance	Xiang <i>et al.</i> , 2007
	<i>OsCIPK12</i> (rice calcineurin B-like protein-interacting protein kinase 12)	Drought tolerance	Xiang <i>et al.</i> , 2007
	<i>OsTPP1</i> (trehalose-6-phosphate phosphatase)	Salinity and cold tolerance	Ge <i>et al.</i> , 2008
	<i>ZFP252</i> (rice TFIIIA-type zinc finger protein)	Drought and salt tolerance	Xu <i>et al.</i> , 2008

tolerance to this condition. Identification and sequencing allow assigning a putative function to a sequence when a significant homology with genes of known function is found. These results are then usually complemented with a proper validation by the use of transgenics. This approach has been especially important in the discovery of several candidate genes in crops in the last decade and, in some cases, it has led to significant improvements in tolerance to stress (Table 3). As an example, the relevance of membrane lipids in tolerance to cold was shown in rice transformed with *Arabidopsis* glycerol-3P-acyltransferase (GPAT) that increased the levels of unsaturated fatty acids in the phosphatidylglycerol by 28% and resulted in a 20% increase in the photosynthetic rates at 17 °C (Yokoi *et al.*, 1998). An example showing the importance of transcription factors in the response to stress was observed in transgenic rice for the transcription factor ABF3 (*Arabidopsis* ABRE-binding factor 3), which showed increased tolerance to drought (Oh *et al.*, 2005).

It is also interesting that different responses are obtained by manipulation of genes within the same family. A good example are calcineurin B-like protein-interacting protein kinases: OsCIPK03, OsCIPK12, and OsCIPK15, whose over-expression in japonica rice Zhonghua, led to specific improved tolerance to cold, drought, and salt stress respectively (Xiang *et al.*, 2007). These results demonstrate the participation of single genes in tolerance to a particular stress. However, it has also been shown that manipulation of single genes can lead to increased tolerance to more than one kind of stress. As an example, the constitutive expression of the transcription factor DREB1A (*Arabidopsis* DRE-binding protein 1) in rice determined increased tolerance to drought and salt stress. Interestingly, when OsDREB1 (rice DRE-binding protein 1) was over-expressed in rice it resulted in increased tolerance for drought, salt and cold stress (Ito *et al.*, 2006). Similar multi-tolerance effects were observed by over-expressing genes such as OsCDPK7, a calcium-dependent protein kinase, which resulted in rice with increased tolerance to cold, salt and drought stress (Saijo *et al.*, 2000). Manipulation of genes with roles other than regulation, such as detoxification, protection and osmotic regulation, has also resulted in increased tolerance to stress in plants (Xu *et al.*, 1996; Capell *et al.*, 2004; Ge *et al.*, 2008; Li *et al.*, 2008; Wang *et al.*, 2008). Targeting effector, rather than regulatory genes, may result in fewer side effects considering the unwanted activation of responsive genes involved in other metabolic pathways.

Despite similarities among different plants, it must not be forgotten that species such as wheat and barley, with far less characterized genomes compared to model plants, may offer unique and interesting features. Their high level of abiotic tolerance and diversity may provide

important resources for validation of candidate genes and accelerate important breeding programs (Langridge *et al.*, 2006). Performance in the field of these species suggests that greater tolerance to abiotic stress is still achievable for other crops if proper research is conducted and should stimulate the exploration of new technologies and alliances between scientists and farmers.

CONCLUSIONS

Gene expression profiling constitutes an exciting tool to unveil mechanisms involved in the response of plants to environmental stress. Its application in crop research is just starting as technologies are becoming more accessible and cost-effective and are expected to fuel huge advances in agriculture in the coming decades. Currently, the importance of biotechnology is being acknowledged by breeding programs around the world and is resulting in the development of new techniques and approaches to increase crop tolerance to stress. Whether Chile will continue to increase its share in the food market worldwide will depend on its ability to develop sustainable and cutting-edge crop research in the future.

RESUMEN

Análisis de la expresión génica: Una forma de estudiar

la tolerancia a estreses abióticos en cultivos. Las regiones agrícolas están cada vez más expuestas a condiciones de frío y sequía, algo especialmente importante en países con opciones limitadas de manejo de cultivos como Chile. Si la creciente demanda por alimento ha de ser cubierta, es necesaria la compleja tarea del desarrollo de nuevos cultivares con mejores rendimientos bajo condiciones de estrés. El desarrollo de la fisiología vegetal y la genómica en los últimos 20 años ha permitido entender mejor cómo las plantas responden al estrés y los mecanismos responsables de los distintos rangos de tolerancia observados en la naturaleza. En esta revisión, se discuten las técnicas más usadas actualmente en análisis de expresión génica y algunas variables que deben ser consideradas en el diseño experimental tales como el estado de desarrollo de la planta y la intensidad, duración e interacción de distintos tipos de estrés, además de la elección de técnicas apropiadas de acuerdo a la información disponible del genoma del cultivo y el objetivo final de la investigación. Todos estos puntos son fundamentales para facilitar la transición desde la genómica a aplicaciones prácticas en el aumento de la tolerancia al estrés de los cultivos. En este sentido, el rápido desarrollo de nuevas técnicas para estudiar la expresión de genes a menor costo determinará una nueva revolución en la investigación de cultivos en

las próximas décadas. En este sentido, Chile necesita estar preparado en esta área para continuar su desarrollo como un importante productor de alimentos a nivel mundial.

Palabras clave: frío, sequía, cultivos, expresión génica.

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USE OF ENZYMATIC BIOSENSORS AS QUALITY INDICES: A SYNOPSIS OF PRESENT AND FUTURE TRENDS IN THE FOOD INDUSTRY

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ABSTRACT

Biosensors are an important alternative in the food industry to ensure the quality and safety of products and process controls with effective, fast and economical methods. Their technology is based on a specific biological recognition element in combination with a transducer for signal processing. The use of enzymatic biosensor technology in food processing, quality control and on-line processes is promising compared to conventional analytical techniques, as it offers great advantages due to size, cost, specificity, fast response, precision and sensitivity. This article reviews the development and use of some enzyme biosensors in the food industry, describes the most important application areas and analyzes the current situation and future possibilities. In conclusion, enzymatic biosensors are a tool with broad application in the development of quality systems, risk analysis and critical control points, and the extent of their use in the food industry is still largely limited by the short lifetime of biosensors, in response to which the use of thermophilic enzymes has been proposed.

Key words: biosensors, enzymes, analysis in food, safety, quality, control processes.

INTRODUCTION

In recent decades increased knowledge about the biological capacity of enzymes has made it possible to create a new generation of products and processes. Among these products are notably biosensors, which represent a powerful alternative to conventional analytical technique (Velasco-García and Mottram, 2003). This technology has advanced considerably in recent years, basically because of the creation of devices applied in the area of biomedicine. These advanced technologies have been gradually transferred horizontally to other sectors, such as the environment and the agro-food industry.

A biosensor is defined as a compact device for analysis that incorporates a biological or biomimetic recognition element (nucleic acid, enzyme, anti-body, receptor, tissue, cell) associated with a transduction system that allows for processing the signal produced by the interaction between the recognition element and the analyte. The principle of detection of a biosensor is based on the specific interaction between the analyte of interest and the recognition element. As a result of this specific interaction, changes are produced

in one or several physical-chemical properties (pH, electron transference, heat transference, change of potential or mass, variation of optical properties, etc.). These changes are detected and can be measured by a transducer (Velasco-García and Mottram, 2003). This system transforms the response of the recognition element into an electronic signal indicative of the presence of the analyte under study or proportional to its concentration in the sample.

Biosensors currently represent powerful tools for analysis with numerous applications in the agro-food industry, mainly in biotechnological instruments (Mello and Kubota, 2002). The most important characteristics of these devices to be competitive with other technologies in the agro-food industry are their specificity, high sensitivity, short response time, their capacity to be incorporated into integrated systems, the facility to automate them, their capacity to work in real time, their versatility and low production cost (Rasooly, 2001; Mello and Kubota, 2002; Velasco-García and Mottram, 2003).

In recent years, the number of scientific investigations and reviews on biosensors has been very high, which reflects the considerable interest in the theme. Ironically, there is a lag between the high level of scientific and technological development and the limited use of these devices in the agro-food sector (Velasco-García and Mottram, 2003) basically because of structural characteristics of the sector, such as legislation, methodological inertia, absorption capacity and environmental factors. The development of the diverse technologies involved in the design and construction of biosensors has allowed in recent years for

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resolving technical difficulties and personalized design of biosensors that, from the technical point of view, cover practically all needs.

The development of biosensors is described in numerous works, the majority in the areas of clinical, environmental, agricultural and biotechnological applications (Tothill, 2001). Their use in the food sector is convenient to ensure the quality and safety of foods (Luong *et al.*, 1991). The potential uses of biosensors in agriculture and food transformation are numerous and each application has its own requirements in terms of the concentration of analyte to be measured, required output precision, the necessary volume of the sample, time required for the analysis, time required to prepare the biosensor or to reuse it and cleanliness requirements of the system (Velasco-García and Mottram, 2003).

Biosensors have been adapted to detect or measure analytes in "on-line" systems, that is, simultaneous with food processing (Rasooly, 2001). Hazard Analysis and Critical Control Point (HACCP) is generally regarded as the most effective system to ensure food safety. It is highly useful in verifying that processes are under control. The high sensitivity of enzymatic biosensors allows for the detection of microorganisms such as *Escherichia coli*, *Salmonella* sp., *Staphylococcus aureus*, pesticides and herbicides, among others, in hours and/or minutes (Fitzpatrick *et al.*, 2000; Killard and Smyth, 2000).

Innovation and development in the food industry are guided by the central principles: food safety and quality (Mello and Kubota, 2002; Ferreira *et al.*, 2003). The increasing complexity of the food chain requires, among other things, the development of effective traceability systems that guarantee the solidity of all the links. In both cases, the priority is to develop and install control systems such as biosensors that involve molecular methods of detection, analysis and diagnosis that are rapid, highly sensitive and automated tracing for a wide range of agents that threaten food safety.

Given the applicability of biosensors, this article reviews the development and use of some enzymatic biosensors in the food industry, describing the three main areas of application: food safety, food quality and process control, the current situation and future possibilities. As well, there is a brief commentary on the aspects of catalytic biosensors and their classification according to the type of interaction established between the recognition element and the analyte.

Enzymatic biosensors

A catalytic biosensor can be described as a compact analytic device that incorporates a biological sensing element, closely connected to or integrated with a transducer system (Velasco-García and Mottram, 2003). Biological sensors include enzymes or multi-enzymatic

mediums, cellular organelles, complete cells or animal or vegetal tissue (Davis *et al.*, 1995; Mello and Kubota, 2002; Wilson and Gifford, 2005), which are used to detect the presence of any of the substrates that participate in the reaction by detecting the disappearance of a known substrate distinct from the substrate being sought, or by the appearance of a known product; biological sensors are not consumed and can be reused (Gajovic *et al.*, 2000; Wilson and Gifford, 2005).

In enzymatic biosensors a reaction occurs catalyzed by an enzyme in which the union of the substrate is produced in a concrete region of the enzyme termed the active center, which upon forming the products is recovered and can begin a new reaction cycle. The use of enzymes as biological recognition elements was very popular in the first generation of development of biosensors owing to their commercial availability or the facility for isolation and purification of diverse sources (Luong *et al.*, 2008). Subsequently, other advantages were found in using enzymes in recognition biosensors, such as rapid response, high selectivity, the possibility of regeneration and the simplicity involved in constructing the devices (Hall, 2002). In numerous cases multi-enzymatic chains are employed, where the enzyme that generally recognizes the analyte does not act directly on it, but rather interacts with some product derived from it. This technique is often used, for example, with some sugars, where enzymes that are used react with the products of hydrolysis of the same. Among the enzymes that are commercially available, the most often used in biosensors are oxidoreductase, notably among which are glucose oxidase, horseradish peroxidase and alkaline phosphatase (Rogers and Mascini, 1998; Laschi *et al.*, 2000), because they are very stable catalyzing reactions of oxide reduction (Mello and Kubota, 2002).

More than 2000 articles on enzyme-based biosensors were found in the literature and this is because of the need to examine blood glucose (Tothill, 2001; D'Orazio, 2003) and the feasibility of construction of the same. In the majority of the applications, the detection limits are satisfactory or excessive, but the stability of the enzymes and the capacity to maintain enzymatic activity over a long period of time continues to be problematic, which is generally resolved by immobilizing the enzymes (Tothill, 2001; D'Orazio, 2003). Immobilized enzymes offer advantages for application in different types of industrial processes and are adaptable to new engineering designs (Krajewska, 2004), for which it is important to increase the affinity of the enzyme to the substrate, reduce inhibition, increase the optimal pH interval and reduce possible microbial contamination (Arroyo, 1998). Basically, in enzymatic biosensors the enzymes are immobilized in a potentiometric, amperometric, optometric, calorimetric or piezoelectric transducer (Davis *et al.*, 1995).

In some cases electro-active interference caused by endogenous compounds in the sample for analysis becomes significant and needs to be eliminated. Currently, glucose oxidase continues to be the most stable and specific enzyme that can be easily obtained in large quantities (Luong *et al.*, 2008).

On the other hand, there are enzymes that cannot be used because they are not sufficiently stable or because their purification is difficult or too costly, because of which cells from bacteria, fungus, protozoa and higher organisms are being used (Mello and Kubota, 2002). Instead of being purified, these cells are used as biological recognition elements, taking advantage of their diverse multi-cellular enzymatic systems that possess the capacity to metabolize different organic compounds, generating distinct products such as ammonia, carbon dioxide, acids, sugars, vitamins and nitrogenated compounds, among others, which can be detected in the biosensors, and in turn can be used to detect compounds that inhibit microbial respiration as toxic and contaminating substances (D'Souza, 2001). As well, the cells offer the facility to be modified genetically to improve their activity or to produce specific enzymes that do not normally appear.

One of the most important limitations in the use of complete cells is the diffusion of substrates and products by means of the cellular membrane, with which a slower response is obtained in comparison to purified enzyme biosensors, and with less specificity owing to reactions catalyzed by other enzymes present in the cell (D'Souza, 2001). These limitations can be reduced through the permeabilization of the cellular membrane by means of enzymatic processes with lysozyme, papain, chemical processes with detergents and physical processes with freezing and thawing (Mello and Kubota, 2002). These processes permit an increase in porosity of the membrane, allowing for better incorporation of the analyte and impede the escape of the intercellular macromolecular compounds, while the enzymes at the same time allow the co-factors that intervene in the catalytic reactions to leave to the exterior of the cell. Consequently, such treatments provoke cellular inviability, which limits their use to applications that do not require cellular regeneration of co-factors or metabolic respiration, as is the case of glucose oxidase, β -galactosidase, amino acid oxidase and invertase (D'Souza, 2001).

Cells can be immobilized in membranes of cellulose acetate, or be trapped in a matrix, such as agar gel (Patel, 2002; Tatsumi *et al.*, 2006; Setti *et al.*, 2007) in a simpler and more economical way than enzymatic immobilization.

Other systems of catalytic biosensors instead of using complete cells or isolated multi-enzymatic include subcellular organelles or tissue, that contain more specific

complete enzymatic systems, as is the case of thylakoids, complete chloroplasts or mitochondria, which are double-membrane organelles that have enzymatic systems related to obtaining energy. Such organelles are used in the detection of toxic agents such as pesticides, heavy metals or detergents that can inhibit enzymatic systems (D'Souza, 2001). In the same manner, there are determined tissues that according to their physiological function in the organism produce specific enzymes or enzymatic systems, such as leaves, roots, fruits or seeds, in sliced or in homogenized form; such tissue are often associated with electrochemical transducers (Li *et al.*, 2002; Wilson and Gifford, 2005; Mei *et al.*, 2007). For example, for the detection of phenol impregnated in salmon, salmon tissue is used in an electrochemical biosensor of carbon and tyrosinase mixed with electropolymerized pyrrole (Tingry *et al.*, 2006); slices of potato are used for the determination of mono and polyphenols (*Solanum tuberosum* L.) because of their high content of polyphenol oxidase enzymes, together with oxygen electrodes (D'Souza, 2001; Kulys and Vidziunaite, 2003); for the determination of alcohol, homogenized fungus *Agaricus bisporus* is used (Akyilmaz and Dinckaya, 2000; Kulys and Vidziunaite, 2003); for the determination of diamines like putrescine and cadaverine, tissue is used from chemiluminescent plants, based on the enzymatic conversion that takes place in the column of tissue of the plants to produce hydrogen peroxide (Mei *et al.*, 2007), among many others examples.

The application of more than one sensor channel for one or more species in a unit using enzyme-electrode type amperometric sensors has led to the development of several multi-sensors based on principles of amperometric detection (Silber *et al.*, 1994; Miertus *et al.*, 1998). The use of electro-chemical transducers of various channels allows for the construction of biosensors that can simultaneously analyze three or more species and in this manner optimize selectiveness and reliability in comparison to sensors with only one substrate (Glazier *et al.*, 1988).

Potential applications of enzymatic biosensors in the agro-food industry

Applications in food safety. The concept of food safety involves ensuring the production and marketing of harmless food, with this, ensure the health of the consumer. The quantity and types of food additives incorporated into food products are regulated by the legislation of each country, the detection and quantification of which are important to prevent fraud and malpractice by manufacturers, allergies and other adverse effects to determined groups of the population (Zinedine *et al.*, 2007). Because of this, special attention has been given to studying the way to detect the presence of contaminants, such as residues of pesticides, fertilizers,

heavy metals and organic compounds, given that the majority of these have a high level of toxicity. Based on this need biosensors are used to detect xenobiotic substances, substances external to the food product such as additives and pesticides and components of the food itself like toxins of diverse origins (Xavier *et al.*, 2000; Patel, 2002). The traditional methods to identify food contaminants include physicochemical, serological and biological tests, however many of these require large quantities of prepared samples, analysis time and lack sufficient sensitivity and selectivity.

The development of catalytic biosensors in food additive analysis generally employs enzymes as recognition systems. This development is described in several investigations, among these notably are analysis of aspartame with carboxyl esterase, alcohol oxidase, caboxypeptidase, L-aspartase, peptidase, aspartate aminotransferase, glutamate oxidase and α -chymotrypsin (Odaci *et al.*, 2004); analysis of sorbitol with sorbitol dehydrogenase and nicotinamide adenine dinucleotide (NAD⁺) (Saidman *et al.*, 2000); analysis of benzoic acid with tyrosine (Morales *et al.*, 2002) and analysis of sulphites with sulphite oxidase; all developed with a system of amperometric transduction. In some cases interference reactions have been observed that reduce the efficacy of these devices, as occurs with biosensors used to detect sorbitol which also interact with another artificial edulcorant, xylitol; and with devices designed for the determination of benzoic acid by the presence of other antioxidants such as butyl hydroxyanisole (BHA) and propyl gallate (Patel, 2002). Table 1 presents the main biosensors used in the detection of these types of compounds in food and water; among these are devices based on the inhibition of enzymatic activities that incorporate enzymes such as cholinesterases (acetyl and/or butyrylcholinesterases), tyrosinase or alkaline phosphatase; and units in which reactions are catalyzed that affect the analyte of interest, which include hydrolases, reductases, etc. (Nunes *et al.*, 1998; Bachmann *et al.*, 2000; Panfili *et al.*, 2000).

The different pesticides used in food production can accumulate in the fatty tissue of animals, while the excessive use of fertilizers contaminates ground water with nitrates, nitrites and phosphates (Cosnier *et al.*, 1998; Moretto *et al.*, 1998). For the detection of herbicides such as phenyl urea and triazines, which inhibit photosynthesis, biosensors have been designed with membrane receptors of thylakoid and chloroplasts, photosystems and reaction centers; or complete cells such unicellular alga and fenilureas and triazines, for which mainly amperometric and optical transducers have been employed (Patel, 2002).

There are also other substances potentially toxic for humans with a major impact on the environment that can reach the food chain accidentally, such as contaminating residues present in water and soil, among these by-products from diverse industrial processes (dioxins), used as dielectric or hydraulic fluid agents (polychlorinated biphenyls or PCBs) or generated in the burning of fossil fuels or wood (polycyclic aromatic hydrocarbons or PAHs), benzene, toluene and xylene (named BETX) and derived phenolics; immunosensors, enzymatic biosensors and biosensors with complete cells are used for the detection of these organic compounds (Hedenmo *et al.*, 1997; Patel, 2002). Likewise, devices have been designed to determine the levels of heavy metals such as arsenic, cadmium, mercury, lead, among others, in samples of water and soil, which incorporate genetically modified microorganisms and enzymes such as urease, cholinesterase, glucose oxidase, alkaline phosphatase, ascorbate oxidase and peroxidase (Tsai *et al.*, 2003), the transduction systems in these devices are notably electrochemical and optical, as indicated in Table 1.

On the other hand, foods can naturally present anti-nutritional compounds that can generate disorders in the consumer, given that they hinder absorption and metabolize distinct nutrients causing a deficiency of the same. Table 2 presents some examples of biosensors used in the detection of anti-nutrients.

Applications in food quality. The term food quality is related to nutritional value, acceptability and safety. The latter was analyzed in the previous section and the others are evaluated in function of parameters such as freshness, appearance, flavor, texture and chemical (Vadiumbal and Jayas, 2007). The composition of the foods allows for characterization and verification if the food contains elements to enrich the food such as vitamins and/or minerals. To evaluate food composition distinct biosensors have been developed, which are described in Table 3.

Various food labeling regulations recognize the importance of determining freshness, establishing guidelines for the use of the term "fresh" in relation to food (FSA, 2004). One way to determine freshness is through evaluation of the composition of products such as meats, fish, fruits and vegetables, given that during periods of storage compounds can be synthesized that produce abnormal odors and flavors and are prejudicial to the health of the consumer. Table 4 lists the biosensors developed to evaluate the freshness and useful life of foods.

Some of the most important problems that affect food freshness, and with it food quality, are exposure time in an inadequate environment, incorrect design of the food packaging, inadequate management of temperatures

Table 1. Most important biosensors used in the detection of pesticides, fertilizers and other pollutants.

Analyte	Type of interaction	Recognition biocatalyzer	Transduction system	References
Pesticides				
Parathion	Biocatalytic	Parathion hydrolase	Amperometric	Velasco-García y Mottram, 2003; Parellada <i>et al.</i> , 1998
Propoxur and carbaryl	Biocatalytic	Acetyl cholinesterase	Fiber optic	Nunes <i>et al.</i> , 1998; Xavier <i>et al.</i> , 2000
Diazinon and dichlorvos	Biocatalytic	Tyrosinase	Amperometric	Pérez Pita <i>et al.</i> , 1997; Mello y Kubota, 2002
Paraoxon	Biocatalytic	Alkaline phosphatase	Optical	Cosnier <i>et al.</i> , 1998; Mello and Kubota, 2002; Patel, 2002
Fertilizers				
Nitrate	Biocatalytic	Nitrate reductase	Amperometric	Moretto <i>et al.</i> , 1998
Nitrite	Biocatalytic	Nitrite reductase	Optical	Moretto <i>et al.</i> , 1998
Phosphate	Biocatalytic	Polyphenol oxidase and alkaline phosphatase, phosphorylase A, phosphoglucomutase and glucose-6-phosphate dehydrogenase	Amperometric	Cosnier <i>et al.</i> , 1998
Heavy metals				
Copper and mercury	Biocatalytic	<i>Spirulina subsalsa</i>	Amperometric	Tsai, 2003; Velasco-García and Mottram, 2003
Copper	Biocatalytic	Recombinant <i>Saccharomyces cerevisiae</i>	Amperometric	Tsai, 2003; Velasco-García and Mottram, 2003
Cadmium and lead	Biocatalytic	<i>Staphylococcus aureus</i> or Recombinant <i>Bacillus subtilis</i>	Optical	Tsai, 2003; Velasco-García and Mottram, 2003
Arsenic, cadmium and bismuth	Biocatalytic	Cholinesterase	Electrochemical	Tsai, 2003; Velasco-García and Mottram, 2003
Cadmium, copper, chrome, nickel, zinc	Biocatalytic	Ureasa	Optical	Tsai, 2003; Velasco-García and Mottram, 2003
Copper and mercury	Biocatalytic	Glucose oxidase	Amperometric	Tsai, 2003; Velasco-García and Mottram, 2003

and the level of oxygen during the handling of fruit and vegetables in modified atmospheres, among many others. Because of this, experimental use has been made of commercial biosensors that use immobilized enzymes like alcohol oxidase and alcohol peroxidase and a chromogene, in which alcohol oxidase catalyzes the oxidation of ethanol in acetaldehyde and H₂O₂ in the presence of O₂, and the peroxidase catalyzes the oxidation of the chromogene, causing a change in color. Smyth *et al.* (1999), measuring with biosensors, ethanol accumulation in lettuce (*Lactuca sativa* L.), cauliflower (*Brassica oleracea* var. *botrytis*),

broccoli (*Brassica oleracea* var. *italica*) and cabbage (*Brassica oleracea* var. *capitata*) lightly processed and packed in a modified atmosphere, detected lesions due to low concentration of O₂ and obtained a response from the biosensor that was very similar to that obtained by gas chromatography, which is costly and requires technical experts. This biosensor can also be used monitor ethanol formation during apple storage in a controlled atmosphere, the development of putrefaction in tubercles like potatoes or for any other application where ethanol accumulation can be associated with quality loss. Likewise, research has

Table 2. Some of the most commonly used biosensors in antinutrient detection.

Analyte	Type of interaction	Recognition system	Transduction system	References
Antinutrients				
Oxalate (spinaches, tea, strawberries)	Biocatalytic	Oxalate oxidase	Amperometric	Milardovic <i>et al.</i> , 2000
Amygdalin (bitter almonds)	Biocatalytic	β -glucosidase	Amperometric	Ohashi and Karube, 1993
			Potentiometric	
Glucoalcaloides	Biocatalytic	Cholinesterase	Potentiometric	Ohashi and Karube, 1993

been conducted that analyzes the content of some organic acids and sugars as indicators of fruit and vegetable maturity (Ángeles and Cañizares, 2004).

There are multiple compounds that give rise to disagreeable flavors and aromas that can be detected with biosensors, as in the case of 2,4,6-trichloroanisole in wine (Moore *et al.*, 2003), which is related to wine bottle corks, whose presence causes significant losses to the wine industry. In other cases, the level of freshness of fish has been detected through a hydrogen peroxide electrode based on the xanthine oxidase enzyme (Volpe and Mascini, 1996). Biosensors can also detect indicators of processes, such as lactulose, disaccharide, which is formed in the thermal treatment of milk allows for distinguishing between milk that has been submitted to a UHT treatment (ultra high temperature) and milk sterilized in the container.

Applications in process control. Currently, thanks to biosensor technology it is possible to determine and quantify on-line diverse compounds of importance in process control, such as sugars, alcohols, and amino acids, among others.

Sugars are limiting factors in fermentative processes given that low concentrations reduce the productivity of the bioreactor. Because of this, numerous investigations have been undertaken, among which notably are those on the use of amperometric biosensors to analyze glucose with glucose oxidase in fruit juices (Ángeles and Cañizares, 2004); lactose with β -galactosidase and glucose oxidase; and lactulose with β -galactosidase and fructose dehydrogenase, which implies an excessive thermal treatment of milk during pasteurization (Campás *et al.*, 2002). In relation to alcohols and principally ethanol, enzymatic reactions are inhibited when alcohol content exceeds 14%; analysis has been advanced mainly with the alcohol dehydrogenase enzyme *Gluconobacter oxydans* with amperometric biosensors; likewise, with fermentation, the proportion of glycerol should be maintained at 1:10 in relation to total alcohol, the analysis of glycerol has been developed with glycerokinase and glycerol-3-phosphate oxidase in amperometric biosensors to monitor fermentative processes (Niculescu *et al.*, 2003). On the other hand, aminoacids like lysine,

obtained by fermentation and employed as animal feed supplements, has been controlled by the lysine oxidase enzyme. Similarly, lactic acid used to control acidity and the formation of crusts on cheese, have been examined with lactate oxidase in amperometric biosensors. These biosensors can be integrated into the system of Hazard Analysis and Critical Control Points (HACCP) to verify that processes are being carried out correctly.

Despite the broad applicability of biosensors, the use of the technology in the control of processes is limited for several reasons: the short life of enzymatic biosensors, the need to calibrate them with certain frequency, the lack of reliable response to different concentrations or with variable conditions in the medium, among others (Ferreira *et al.*, 2003). The prototype tests in real samples have critical stages such as immobilization of the biocomponent during the construction of the device and preparation of the sample for analysis. Biosensors require mild temperature and pH conditions to keep the biological element active (Gibson, 1999; Wilson and Gifford, 2005). Consequently, in some cases a previous treatment of the sample is recommended to eliminate interfering species such as ascorbic acid, tyrosine and others. Procedures are conducted that include neutralization, dilution or extraction when the food is acidic or hydrophobic. The correction methods to reduce the duration of food analysis include acidic or alkaline hydrolysis, microwave digestion, extraction of supercritical fluids, evaporation and filtration (Deng and Dong, 1996; Marconi *et al.*, 1996; Kotsira and Clonis, 1998; Panfili *et al.*, 2000). Likewise, specific sensors have been developed for the determination of glucose, lactate, glutamate, pyruvate, choline and acetylcholine through monitoring of nitric oxide, Na^+ , K^+ , Ca^{2+} , and dopamine (Zhang and Wilson, 1998; Wilson and Gifford, 2005).

CONCLUSIONS

The food industry is benefitting from major advances in the development of enzymatic biosensors with different transduction systems that can be applied in the areas of food safety, quality and process control; studies are focused mainly on determining composition, contamination of primary materials and processed foods.

Table 3. Most important biosensors applied to evaluate food quality.

Analyte	Matrix	Recognition enzyme	Transduction system	References
Glucose	Grape juice, wine, juice, honey, milk and yogurt	Glucose oxidase	Amperometric	Centonze <i>et al.</i> , 1997; Ángeles y Cañizares, 2004
Fructose	Juice, honey, milk, gelatin and artificial edulcorants	Fructose dehydrogenase, D-fructose 5-dehydrogenase	Amperometric	Bassi <i>et al.</i> , 1998; Palmisano <i>et al.</i> , 2000
Lactose	Milk	β -Galactosidase	Amperometric	Marconi, 1996; Palmisano <i>et al.</i> , 2000
Lactate	Cider and wine	Transaminase and L-lactate dehydrogenase	Amperometric	Silber <i>et al.</i> , 1994; Ramanathan <i>et al.</i> , 2001
Lactulose	Milk	Fructose dehydrogenase and β -galactosidase	Amperometric	Sekine and Hall, 1998
L-amino acids	Milk and fruit juices	D-amino acid oxidase	Amperometric	Sarkar <i>et al.</i> , 1999
L-glutamate	Soya sauce and condiments	L-glutamate oxidase	Amperometric	Matsumoto <i>et al.</i> , 1998; Kwong <i>et al.</i> , 2000
L-lysine	Milk, pasta and fermentation samples	Lysine oxidase	Amperometric	Kelly <i>et al.</i> , 2000; Olschewski <i>et al.</i> , 2000
L-malate	Wine, cider and juices	Dehydrogenated malate, others	Amperometric	Miertus <i>et al.</i> , 1998
Ethanol	Beer, wine and other alcoholic drinks	Alcohol oxidase, alcohol dehydrogenase, NaDH oxidase	Amperometric	Katrlık, 1998 ; Miertus <i>et al.</i> , 1998
Glycerol	Wine	Glycerophosphate oxidase and glycerol kinase	Amperometric	Niculescua <i>et al.</i> , 2003
Catechol	Beer	Polyphenol oxidase	Amperometric	Eggins <i>et al.</i> , 1997
Cholesterol	Butter, lard and egg	Cholesterol oxidase and peroxidase	Amperometric	Akyilmaz and Dinckaya, 2000.
Citric acid	Juice and athletic drinks	Citrate lyase	Amperometric	Prodromidis <i>et al.</i> , 1997
Lecithin	Egg yolk, flour and soya sauce	Phospholipase D and choline oxidase	Electrochemical	Mello and Kubota, 2002

In the area of food safety, enzymatic biosensors allow for identifying the presence of highly toxic organic contaminants and the presence of anti-nutritional elements that affect the food chain, either accidentally or by intention. This early detection protects the environment from contaminants and consumers from chronic illnesses and allergies.

Equally, enzymatic biosensors are being used in the food industry to determine the freshness of products given that it is possible to detect enzymes and compounds of aroma and flavor that originate from the senescence stage of products.

Biosensors have proven to be especially useful in the control of fermentative processes in follow-up of the consumption of the substrate by microorganisms, control of acidity and assessing the thermal profile.

While the use of biosensors in the food industry is on a mass scale, there are still obstacles to be overcome, such as the high cost of purifying the enzymes that are used as detecting elements, the low specificity and low response time that are obtained when complete cells or tissue are used, the lack of reliable responses at low concentrations, interference reactions, the need to calibrate the devices and the stability of the enzymes. This last factor is the most limiting for the lifetime of enzymatic biosensors. If these limiting factors can be overcome, it will be possible to develop enzymatic biosensors that are more rapid, versatile, reliable, long lasting and cost-effective.

Cuadro 4. Biosensor used in the evaluation of freshness and self life.

Analyte	Matrix	Recognition enzyme	Transduction system	References
Evaluation of spoilage				
Polyphenols	Olive oil	Tyrosinase, laccase	Amperometric	Campanella <i>et al.</i> , 1993; Kuly and Vidziunaite, 2003
Short chain fatty acid	Milk and derivatives	Lipase	Electrochemical	Mello and Kubota, 2002
Freshness index				
Ornithine and amines	Shrimps	Ornithine carbamoyl transferase, nucleoside phosphorylase and xanthine oxidase	Amperometric	Mello and Kubota, 2002
Amines	Fish, lobster	Diamine oxidase, Ornithine carbamoyl transferase, nucleoside phosphorylase	Amperometric	Park <i>et al.</i> , 2000; Mello and Kubota, 2002
Biogenic amines	Fish	Amine oxidase and peroxidase	Amperometric	Tombelli and Mascini, 1998
Hypoxanthine	Fish	Xanthine oxidase	Amperometric	Hu <i>et al.</i> , 2000
Lactic acid	Meat	Xanthine oxidase, diamine oxidase Polymide oxidase	Amperometric	Mello and Kubota, 2002
Evaluation of maturity				
Glucose	Fruit	Glucose oxidase	Electrochemical	Ramanathan <i>et al.</i> , 2001
Sucrose	Fruit	Invertase, mutarotase and glucose oxidase	Electrochemical	Mello and Kubota, 2002
Isocitrate	Fruit	Isocitrate dehydrogenase	Potentiometric	Mello & Kubota, 2002

RESUMEN

Uso de biosensores enzimáticos como indicadores de calidad: Una sinopsis del presente y futuro en la industria alimentaria. Los biosensores constituyen una importante alternativa en la industria de alimentos para garantizar la calidad e inocuidad de los productos y controlar los procesos con métodos eficaces, rápidos y económicos; su tecnología está basada en un elemento de reconocimiento biológico específico en combinación con un transductor para el procesamiento de la señal. El uso de técnicas de biosensores enzimáticos en procesamiento de alimentos, control de calidad y de procesos "on line", es prometedor frente a las técnicas analíticas convencionales, ya que ofrecen grandes ventajas debido a su tamaño, costo,

especificidad, respuesta rápida, precisión y sensibilidad. En este artículo se revisa el desarrollo y uso de algunos biosensores enzimáticos en la industria alimentaria, se describen las áreas de aplicación más importantes y se analiza su situación actual y posibilidades futuras. En conclusión, los biosensores enzimáticos son una herramienta de gran aplicabilidad en el desarrollo de sistemas de calidad como el análisis de riesgos y puntos críticos de control, y que la masificación de su uso en la industria alimentaria se ve aún limitada principalmente por el tiempo de vida útil de los biosensores, para lo cual se propone el uso de enzimas termofílicas.

Palabras clave: biosensores, enzimas, análisis en alimentos, seguridad, calidad, control de procesos.

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EFFECT OF PHEROMONE TRAP DENSITY ON MASS TRAPPING OF MALE POTATO TUBER MOTH *Phthorimaea operculella* (ZELLER) (LEPIDOPTERA: GELECHIIDAE), AND LEVEL OF DAMAGE ON POTATO TUBERS

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ABSTRACT

Potato tuber moth (PTM), *Phthorimaea operculella* (Zeller), is one of the pests that cause the most damage to potatoes (*Solanum tuberosum* L.) in both field crops and storage, especially in regions where summers are hot and dry. Larvae develop in the foliage and tubers of potatoes and cause direct losses of edible product. The use of synthetic pheromones that interfere with insect mating for pest control has been widely demonstrated in numerous Lepidoptera and other insect species. An experiment was carried out during the 2004-2005 season in Valle del Elqui, Coquimbo Region, Chile, to evaluate the effectiveness of different pheromone trap densities to capture *P. operculella* males for future development of a mass trapping technique, and a subsequent decrease in insect reproduction. The study evaluated densities of 10, 20, and 40 traps ha⁻¹, baited with 0.2 mg of PTM sexual pheromone, and water-detergent for captures. Results indicated that larger numbers of male PTM were captured per trap with densities of 20 and 40 traps per hectare, resulting in a significant reduction ($P < 0.05$) of tuber damage in these treatments compared with the control which used conventional chemical insecticide sprays.

Key words: potato tuber moth, *Phthorimaea operculella*, mass trapping, pheromone.

INTRODUCTION

The potato tuber moth is a pest which economically affects potato crops, mainly in regions where the climate is hot and dry. The larvae cause direct damage to the tubers by infesting them underground and control using chemical insecticide sprays is difficult with uncertain results.

The use of synthetic sexual pheromones to interfere with reproduction offers a non-traditional way to manage pest control that does not use insecticides. Sexual pheromones are species-specific and highly selective, and since they are not toxic and do not represent health risks to humans and animals, they are valuable tools in integrated pest control management.

The use of pheromone traps for mass trapping is an insect control method that has been sufficiently

researched (El-Sayed *et al.*, 2006). It interferes with insect mating, reducing the future larvae population and subsequent damage. In order to improve the effectiveness of the captures and make the traps a more reliable tool in management programs, it is necessary to determine the factors that affect their efficiency (Athanasios *et al.*, 2002; 2003a; 2003b; 2004; 2005; 2007). Traps can also be used with the degree-day calculation method for decision-making on the application of insecticides in pest control (Kumral *et al.*, 2005).

The *P. operculella* pheromone has been studied since 1969 by Adeesan *et al.* who discovered that it was released from a gland located just before the last abdominal segment of the female. Afterwards, research by Fouda *et al.* (1975), Roelofs *et al.* (1975), Persoons *et al.* (1976), Yamoaka *et al.* (1976), Bacon *et al.* (1976), Voerman *et al.* (1977), Voerman and Rothschild (1978), identified, isolated, and synthesized the two main pheromone compounds, finding that combining them attracted the male moths more than each compound taken separately. Bacon *et al.* (1976) and Raman (1982; 1984) also evaluated the most effective rates of these two compounds in the mixture preparation, as well as some techniques to use these compounds in field and storage traps.

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This pheromone is currently available on the market and is related to the mixture of the *trans*-4, *cis*-7-tridecadienil-1-ol-acetate and *trans*-4, *cis*-7, *cis*-10 tridecatrienil-1-ol-acetate compounds in a ratio of 1:1.5. This mixture is generally commercialized as a rubber device in which the mixture is impregnated in 1 mg doses.

Though there are some control studies of mass trapping and mating disruption of *P. operculella* (Raman, 1982; 1984; Salas *et al.*, 1985; Ortu and Floris, 1989), these techniques are not used and this pest is mainly controlled with broad-spectrum pesticides.

On the other hand, since there are many factors that can affect the effectiveness of these control methods, it is important to conduct studies that confirm the effectiveness of pheromone use as a control technique for specific crop conditions where the moth constitutes an economically important pest. This explains why the objective of this study was to evaluate the effect of different trap densities on the number of male *P. operculella* captured and the effect of these captures on the reduction of damage caused by larvae in tubers produced under agro-ecologic conditions in the coastal zone of the Coquimbo Region, Chile.

MATERIALS AND METHODS

During the 2004-2005 season, a massive trapping trial of male potato tuber moth (PTM) was conducted using pheromones in different densities of traps per area. The trial was carried out in El Romero sector (29°53' S; 71°07' W), La Serena, Coquimbo Region, Chile, using a randomized complete block design with four replications. The area of the experimental plots was 4000 m².

Sowing took place on 30 October 2004 in an area of approximately 13 ha. Nine hectares were sown with certified daughter seed of the Asterix variety, whereas the remaining hectares were sown with Cardinal potato seed.

The trial was fertilized with N, P₂O₅, and K₂O in doses of 150, 120, and 60 kg ha⁻¹, respectively. Mancozeb 1.6 kg a.i. ha⁻¹ (Mancozeb 80% WP) fungicide was applied during cultivation to control late blight (*Phytophthora infestans*). Linuron 1 kg a.i. ha⁻¹ (Linurex 50 WP) was initially applied to control weeds, but these were later controlled manually. Furrows were irrigated every 7 days. Harvest took place between 31 January and 4 February 2005.

Three trap densities were evaluated with the sexual pheromone of *P. operculella*: *trans*-4, *cis*-7-tridecadienil-1-ol-acetate and *trans*-4, *cis*-7, *cis*-10 tridecatrienil-1-ol-acetate in a ratio of 1:1.5, and 0.2 mg dose per trap in accordance with the results obtained by Larraín *et al.* (2007). The trap densities evaluated were 10, 20, and 40 traps ha⁻¹ in plots untreated with insecticides. Treatments were distributed within each plot, with 35 m between

plots and a minimum of 5 m between traps (40 traps ha⁻¹) (Figure 1). These distances were selected in order to avoid the effect between treatments with distinct trap densities, and considering the results of Cameron *et al.* (2002), who studied the activity of *P. operculella* with entomologic nets and pheromones, finding that adult activity declined at distances of 20 and 40 m from a release point.

The number of traps used in the trial area corresponding to the 10, 20, and 40 trap ha⁻¹ densities were 4, 8, and 16 treatments, respectively, with a total of 112 traps in the trial. The traps were all assembled in the same way as established by Larraín *et al.* (2007), using 5 L capacity plastic drums containing 2 L of water with 0.2% detergent. The dispenser with 0.2 mg of pheromone per trap was supplied by Agrisense-BCS (Pontypridd, South Wales, UK).

Evaluations

All traps were set up on 6 December 2004. Starting on that date and until harvest, a weekly count checked the number of male moths captured in all the traps of each plot. At that moment, an 80 kg sample of tubers from a 25-30 m² area was taken. This sample was taken from the center of each 4000 m² plot. Four plots of the same area were selected as controls, chosen randomly in the same field, at a distance beyond the influence of the pheromone. Due to the high insect pressure during the season, these plots were managed by the farmer who had to spread eight applications using a permethrin (Pounce[®]) and methamidophos (MTD 600 SL, ANASAC, Chile) mixture in doses of 1 L and 200 cm³ ha⁻¹, respectively.

The damage caused by *P. operculella* was evaluated in all tubers of the samples. The number of undamaged tubers was counted and the percentage of moth damage was calculated in each treatment.

Data from the number of captures and the percentage of damaged tubers was analyzed using a variance analysis (ANOVA) with 5% probability and the means were compared by using a multiple comparison test with a Least Significant Difference (LSD). The statistical software used was SAS 8.0 for Windows.

RESULTS AND DISCUSSION

Effect of trap density on captures

P. operculella captures increased as the trap density increased (Table 1). Densities of 20 and 40 traps ha⁻¹ captured a significantly higher number ($P < 0.05$) of males than the 10 traps ha⁻¹ density. However, between the higher density traps (20 and 40 traps ha⁻¹), the mean of males captured was not significantly different ($P \geq 0.05$). The mean of males captured per hectare indicates that approximately

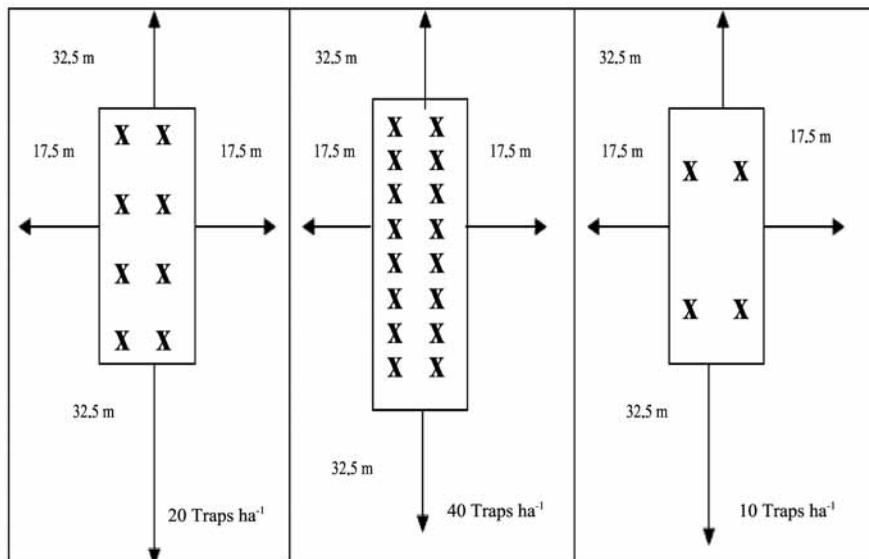


Figure 1. Distribution of treatments in trap density trial. El Romero 2004-2005.

108 588 and 90 173 males ha⁻¹ were captured with densities of 40 and 20 traps ha⁻¹, respectively (Table 1). These results were higher than those obtained by Raman (1988) in Lima, Peru, who captured 92 000 male *P. operculella*, but in 112 d with a density of 42 traps ha⁻¹.

These differences could be due to various factors, such as the pressure of the male moth population in relation to its location, and trap characteristics (type, size, and pheromone dose per trap). Raman (1988) used a standard capsule from Centro Internacional de la Papa (CIP) (International Potato Center) with a 1 mg dose that was less attractive than the 0.2 mg used in this trial according to the results of Larrain *et al.* (2007).

On the other hand, Ortu and Floris (1989) increased to 84 traps ha⁻¹, and observed a drastic reduction in captures, indicating that with this density, the sexual confusion of *P. operculella* could start to be effective.

Tuber damage by moths

Damage caused by *P. operculella* larvae in the tubers was significantly higher ($P < 0.05$) with the control using insecticide treatment (51% damaged tubers) than in the

plots with different densities of pheromone traps evaluated (Table 2).

In accordance with the results obtained by Araya *et al.* (2000), the potato tuber moths in La Serena, Chile, were less susceptible to methamidophos than populations evaluated in other regions in this country. However, no resistance was noted, so failure of the control could be due to the difficulty of insecticides to reach the insects once they are underground or inside the tubers, and also probably due to a higher mortality of the natural enemies of *P. operculella* caused by the synergetic effect of two broad-spectrum action insecticides.

On the other hand, in plots treated with pheromones, the highest rate of tuber damage (30%) was obtained with the lowest density of 10 traps ha⁻¹, and less captures. It is possible to emphasize that the damage was similar to that obtained in plots with 40 traps per hectare (25%), where captures almost doubled compared to 10 traps ha⁻¹ (Table 2). This aspect should be clarified with new studies where more data will be obtained to allow the analysis and establishment of a relation between the male captures and the tuber damage caused by the moth.

Raman (1988) observed minor damage that reached 19% with lower captures and a similar trap density, though factors such as varietal susceptibility, temperatures during the months of capture, pheromone dose per trap, and type of trap do not allow the comparison of these results.

In plots with moderate density of 20 traps ha⁻¹ with pheromones, minor damage was found with 18.3% of the tubers affected and significant differences ($P < 0.05$) compared to the lesser density evaluated in 10 traps ha⁻¹ (Table 2).

Table 1. Mean of *Phthorimaea operculella* males captured in the 2004-2005 crop season.

Treatment (N° traps ha ⁻¹)	Number of males captured (64 days)
40	108 588a
20	90 173a
10	54 864b
Coefficient of variation, %	17.5

Previous results indicate that the control of *P. operculella* with the insecticides most used in the Valle del Elqui, Chile, is inadequate since the number of tubers damaged was higher than 50% with eight applications (Table 2). This entails a high economic cost for the farmer and generates environmental damage since insecticides are applied excessively, for example, methamidophos, which is one of the most toxic of the organic phosphate group and classified as an extremely dangerous toxicological category.

This is the first study that analyzes the potential use of mass trapping with sexual pheromones as a direct control method of *P. operculella* in Chile. The results reflect that the use of this technique constitutes a tool which can significantly reduce the number of males with a subsequent decrease in potato tuber moth larvae, resulting in a significantly lower tuber damage rate in the area within pheromone influence.

Table 2. Mean percentage of tubers damaged by *Phthorimaea operculella* in different pheromone trap densities and control (pesticide management).

Treatment (traps ha ⁻¹)	Mean of tuber damage (%)
Control	51a
10	30b
20	18c
40	25bc
Coefficient of variation, %	19.4

CONCLUSIONS

Twenty traps per hectare appear to be the most effective and convenient trap density to use in a potato tuber moth integrated management program. No significant difference was shown in male capture and tuber damage between this density and the higher trap density trials, therefore the lower density trap pattern proved to be as efficient and more economical.

RESUMEN

Efecto de la densidad de trampas de feromona en la captura masiva de machos de polilla de la papa, *Phthorimaea operculella* (Zeller) (Lepidoptera: Gelechiidae), y en el nivel de daño a los tubérculos.

La polilla de la papa, *Phthorimaea operculella* (Zeller), es una de las plagas que causan mayor daño a la papa (*Solanum tuberosum* L.), tanto a los cultivos en campo como a los tubérculos almacenados, especialmente en zonas de climas cálidos y secos. Las larvas de este insecto se desarrollan en el follaje y tubérculos de papa causando pérdidas directas del producto a comercializar. La

utilización de feromonas sintéticas, como una herramienta que interfiere con el apareamiento, ha sido ampliamente demostrada en innumerables especies de polillas y otros insectos. Con el fin de evaluar la efectividad de diferentes densidades de trampas de feromona en la captura de machos de *P. operculella*, para su futura utilización como técnica de trampeo masivo y consecuente disminución de la reproducción del insecto, se realizó un estudio durante la temporada 2004-2005, en el Valle del Elqui, Región de Coquimbo, Chile. Se evaluaron densidades de 10, 20 y 40 trampas ha⁻¹ con una carga de 0,2 mg de feromona por trampa, utilizando trampas de agua con detergente para las capturas. Los resultados indican que la mayor captura de machos de polilla de la papa se obtiene con densidades de 20 y 40 trampas ha⁻¹, encontrándose también una reducción significativa ($P < 0,05$) del daño en tubérculos en estos tratamientos comparados con el testigo convencional con aspersiones de insecticidas.

Palabras clave: papa, plagas, polilla de la papa, *Phthorimaea operculella*, capturas masivas, feromonas.

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RHIZOGENIC INDUCTION IN ADULT *Juglans regia* L. cv. SERR TISSUE INDUCED BY INDOLE BUTYRIC ACID AND *Agrobacterium rhizogenes*

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ABSTRACT

The *in vitro* introduction of adult walnut (*Juglans regia* L.) tissue represents an opportunity to clone elite genotypes whose selection occurs in advanced ontogenic states. With the purpose of developing a protocol to allow mass propagation of valuable genotypes from adult material, a comparison was made between two root induction systems of walnut microshoots of the fourth subculture of adult walnut tissue of an *in vitro* introduction program previously reinvigorated through traditional grafting. Rhizogenic induction by indole-3-butyric acid (IBA) and *Agrobacterium rhizogenes* was used. The rhizogenic process was analyzed in two phases for both auxinic (T1: 3 mg L⁻¹ IBA; T2: 5 mg L⁻¹ IBA) and *A. rhizogenes* inductions (T3: A-477; T4: A-478). The first phase of root induction was during 3 days in the dark while the second phase, root manifestation, was 27 days. Rooting percentage was evaluated and the induced root systems characterized (number, length, diameter, and root insertion zone) in all the procedures. The best rooting results were obtained in T2, although the response obtained with *A. rhizogenes* didn't differ from the T1 response. This appears to be an increasingly interesting methodology for adventitious rhizogenesis in this species.

Key words: rooting, microshoots, adult material, *Agrobacterium rhizogenes*.

INTRODUCTION

The application of walnut regeneration methods by means of *in vitro* culture of embryos has allowed overcoming the barriers for the large scale production of crops, such as low percentage of seed germination and long propagation cycles. In the first case, between two and three months of stratification are required, whereas the propagation cycles are related to obtaining appropriate size patterns for grafting and the development of commercial specimens in a period of two to three years. With respect to the formation of microplants by means of *in vitro* culture, plants with intact roots, shoots, and leaves have been obtained in distinct culture mediums (Leslie and McGranahan, 1992; Driver y Kuniyuki, 1994; Sánchez-Olate *et al.*, 1997; Fernández *et al.*, 2000), finding that roots were more robust and developed than leaves (Kaur *et al.*, 2006). However, there is no record of the utilization of this technique in adult *J. regia* material which is of vital importance for the development of a massive propagation program, taking into consideration that it is a recalcitrant species (Preece *et al.*, 1989; Leslie and McGranahan, 1992; Caboni *et al.*, 1996; Rodríguez

et al., 2005) with a reduced morphogenetic capacity in the adult phase due to a complex metabolic and tissular system (Sánchez-Olate *et al.*, 2002). These characteristics make it necessary to search for reinvigoration techniques appropriate to this species, such as the severe pruning applied to *Corylus avellana* L. (Sánchez-Olate *et al.*, 2004) and *Pinus radiata* D. Don (Materán *et al.*, 2008).

In vitro reinvigoration techniques have been applied which allow the establishment of a propagation system on a larger scale from adult material selected for productivity. However, the presence of various phenol endogenous compounds, including allelopathic naphthoquinone called juglone, which interfere with cell growth (Fernández *et al.*, 2000) have made it difficult to obtain successful results. It has been suggested that the presence of polyamines, endogenous juglone, or the continuity of sclerenchymatic cylinders contained in the phloem of plant material inhibits root formation (Günes, 1999). This would explain the fact that only high rhizogenic rates have been obtained from material of embryonic origin (Leslie and McGranahan, 1992). This fact restricts the advantages that elite genotype cloning represents where selection occurs in advanced ontogenic states with a high complexity at both the metabolic and tissular levels (Sánchez-Olate *et al.*, 2002).

Microshoots of micro-propagated *in vitro* *J. regia* from isolated embryos have been rooted using auxins such as naphthalene acetic acid (NAA) and indole butyric

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acid (IBA) (Ripetti *et al.*, 1994), genetic transformation with *Agrobacterium tumefaciens* (McGranahan *et al.*, 1988), and inoculation with *A. rhizogenes* (Caboni *et al.*, 1996). The latter permits the induction of adventitious roots in the infection zone due to the transfer of genetic information of a portion of (T-DNA) of Ri (Root inducing) plasmid from the bacteria to the plant genome (Strobel and Nachmias, 1988). Caboni *et al.* (1996) used this method to achieve rooting rates between 52 and 68% in embryonic *J. regia* cv. Sorrento microshoots that were successfully transferred *ex vitro*. At this moment, no results using adult material have been reported.

The results obtained with *Agrobacterium* have been related to a synergy between the auxinic (IBA) concentration and the *A. rhizogenes* infection which is expressed by a possible response of living plant cells contiguous to the dead cells infected with the bacteria. The living cells would transmit diffuse signals to other healthy cells that are capable of initiating the rhizogenesis process (Falasca *et al.*, 2000). According to Vahdati *et al.* (2002), the *rol* genes derived from the T-DNA of *A. rhizogenes* are involved in changing the following characteristics in the transformed plants: *rol A*: wrinkled leaves, condensed inflorescences, increment in the size of the stigma and large flowers; *rol B*: increments the rooting potential as a result of increasing sensitivity to the tissue auxins, alters the morphology, and increases flower size; *rol C*: reduces internode length, produces flowering abnormalities, and increments ramification; *rol D*: causes dwarfism and early blooming. However, in transgenic 5-year old walnut trees transformed by the *rol ABC* genes, no differences were found in the growth habit of shoots and roots (Vahdati *et al.*, 2002).

The effect of indole butyric acid and two wild strains of *Agrobacterium rhizogenes* on rhizogenic induction in *Juglans regia* microshoots was studied since the aim was to develop a protocol allowing massive propagation from adult material.

MATERIALS AND METHODS

Plant material

Caulinar portions of adult material were used and obtained from epicormic shoots of reinvigorated material by grafting *J. regia* cv. Serr on *J. nigra*. This was maintained in the fourth subculture on a DKW (Driver and Kuniyuki, 1994) proliferation medium with pH 5.8, supplemented with sucrose (3%), benzylaminopurine (BAP) (1 mg L⁻¹), IBA (0,01 mg L⁻¹), and gelled with agar agar (7 g L⁻¹) in photoperiodic environmental conditions of 16 h, 25 ± 1°C during the day, 22 ± 1°C during the night, 60% relative humidity, and light intensity of 40 μE m⁻²s⁻², as reported by Sánchez-Olate *et al.* (2002).

Rooting assay

The rhizogenic process was analyzed for auxinic induction (T1: 3 mg L⁻¹ AIB; T2: 5 mg L⁻¹ IBA) and *A. rhizogenes* induction (T3: A-477; T4: A478) in two phases. The first phase was root induction in the dark for 3 days, and the second 27-day phase was known as root manifestation.

Rooting induced by IBA

Microshoots, 3 cm long, were handled using the rooting methodology described by Ripetti *et al.* (1994), and tested in two concentrations of exogenous IBA (3 and 5 mg L⁻¹) in an MS (Murashige and Skoog, 1962) medium with 25% (MS¼) macronutrients. An induction phase of 3 days was maintained in the dark at a temperature of 25 ± 1 °C during the day, 22 ± 1 °C during the night, and 60% relative humidity. Once the induction phase was finalized, the microshoots were transferred to the root manifestation phase of 27 days in a 16:8 photoperiod in a DKW (Driver and Kuniyuki, 1994) medium with 25% (DKW¼) macronutrients mixed with vermiculite (220/250 v/v), and solidified with gelrite (Phytigel, Sigma®).

Rooting induced by *A. rhizogenes*

Microshoots, 3 cm long, were inoculated in their basal portion with A-477 and A-478, wild strains of *A. rhizogenes* from the Valencia, Spain collection (Dawson *et al.*, 1990). The reactivation of bacterial growth was carried out with an aliquot in a microbiological beaker and by resuspending it in a 2 mL liquid medium of YMB (Yeast Medium Basal) (Hooykaas *et al.*, 1977), and shaking it during 48 h at 300 rpm and 25-27 °C. Subsequently, 100 μL of aliquots were taken from the initial bacterial suspension to be resuspended in 10 ml capacity tubes containing 2 mL of liquid YMB medium to reapply the initial treatment. Finally, 100 μL of aliquots were cultivated in a solidified medium with agar (8 g L⁻¹) on Petri dishes with 10 mL of medium maintained at 28 °C during 24 h in inverted position to avoid evaporation.

Once the colonies were developed, a 100 mL Erlenmeyer flask containing 20 mL of liquid YMB medium was inoculated with an inoculation loop by shaking it at 300 rpm at 25-27 °C for 48 h. Subsequently, the solution was placed on a sterile Petri dish in order to proceed with the inoculation of the microshoots obtained from the proliferative chains. The basal inoculation of the microshoots was carried out by submerging them during 3 min in a bacterial solution after eliminating the basal axillary buds and cutting them in 1 cm lengths to increase the infection area. The microshoots were immediately placed on sterile filter paper arranged in a laminating flow chamber to dry and afterwards cultivate them in glass containers with 25 mL of MS¼ medium during three days

in a dark chamber at a temperature of 25 ± 1 °C during 16 h, 22 ± 1 °C during 8 h, and 60% relative humidity. At the end of the induction phase (3 days in the dark), the microshoots were cultivated in a DKW¼ medium mixed with vermiculite (200/250 v/v), and solidified with gelrite (2,5 g L⁻¹ of Phytigel, Sigma®) to which 300 µg mL⁻¹ of Cephotaxime (Claforan® 1 g, Roussel Ibérica, S.A.) were added to control bacterial development. The environmental conditions corresponded to a 16 h photoperiod of light during 27 days at a temperature of 25 ± 1 °C during the day, 22 ± 1 °C during the night, and 60% relative humidity.

The experimental design was completely random with four replications. The experimental unit corresponded to a container with four microshoots each measuring 2.5 cm. At the end of the manifestation period the microshoots were extracted and carefully washed to eliminate the substrate adhering to the roots. The treatments were compared by evaluating the percentage of rooting and the induced radicular systems, contrasting them statistically with ANOVA. Significant differences were identified with the Tukey multiple comparison test with 95% probability (Steel and Torrie, 1985).

RESULTS

Results indicated that it is possible to induce adventitious rhizogenesis in microshoots originating from adult material (Table 1) when they are partially reinvigorated by grafting. The rates of rooting and the quality of the resulting radicular system are similar to those obtained in microshoots of embryonic origin (Sánchez-Olate, 1997), but with different responses depending on the inductor used.

As occurs with material of embryonic origin, results showed a close relationship between callogenesis and rhizogenesis. In each treatment, as the percentage of callogenic tissue increased, a smaller number of explants rooted (Table 1). Even with the best rooting results obtained in T2 (Table 1), the response to *A. rhizogenes* (T3

and T4) did not differ from T1. Hence, it could become an increasingly interesting method for adventitious rhizogenesis in this species since observed rhizogenic rates greatly exceeded the results reported by Sierra (2002) in cutting rooting of the same species (12%).

By analyzing the number of roots for each microshoot, T2 generated a significantly greater response with respect to T1, T3, and T4. In the latter two, as occurred in the rooting rate, there were no significant differences (Table 1). On the other hand, major differences occurred in the length of the induced radicular system since the 5 mg L⁻¹ application of IBA produced significantly larger roots than in the other treatments for the same manifestation period (Figure 1a and 1b). This led to two hypotheses: a) that the high auxinic concentration forces an accelerated metabolic route toward the synthesis of rhizogenic tissues, or b) that the high auxinic concentration forces a strong initial pulse, the roots are generated in the first hours of manifestation and develop in length over an extended period of time. This second hypothesis coincided with observations made by Ríos *et al.* (2002) in cotyledonal portions of the same species where it was observed that the rhizogenic induction produced by high auxinic concentrations took place in the first few hours after applying treatment. This resulted in a numerous and longer radicular system with respect to the treatments with lesser concentrations of IBA and greater concentrations of other auxins such as indolacetic acid (IAA) and NAA.

With respect to the induction by *A. rhizogenes*, the results indicate that this induction agent seems to be a real alternative to rooting of this species, not only because of its easy application, but the acceptable rooting rates achieved (Table 1) and the radicular systems obtained (Figure 1c and 1d). Furthermore, the greatest area of influence of this induction agent, manifested by the appearance of roots in the whole area treated, generated their better distribution in relation to its insertion in the caulinar portion.

The differences between the materials obtained in both *A. rhizogenes* strains basically rested on the fact that T3 obtained longer roots (Figure 1c). Perhaps the most

Table 1. Rhizogenic evaluation induced by indole-3-butyric acid (IBA) and *Agrobacterium rhizogenes*.

Treatment	%		N° of roots	mm		Root insertion zone ¹
	Callogenesis	Rhizogenesis		Root length	Root diameter	
T1	66.7b	50.0a	1.8a	09a	1a	B
T2	50.0a	67.5b	3.1b	18b	2b	MB
T3	90.3c	47.7a	2.3a	09a	2b	SMB
T4	81.3c	50.0a	2.0a	05a	2b	SMB

Different letters show significant statistical differences ($\alpha \leq 0, 05$).

¹ Root insertion zone to the shoot. S: superior zone of the portion in contact with the culture medium. M: medium zone. B: basal zone. T1: 3 mg L⁻¹ IBA. T2: 5 mg L⁻¹ IBA. T3: A-477. T4: A-478.

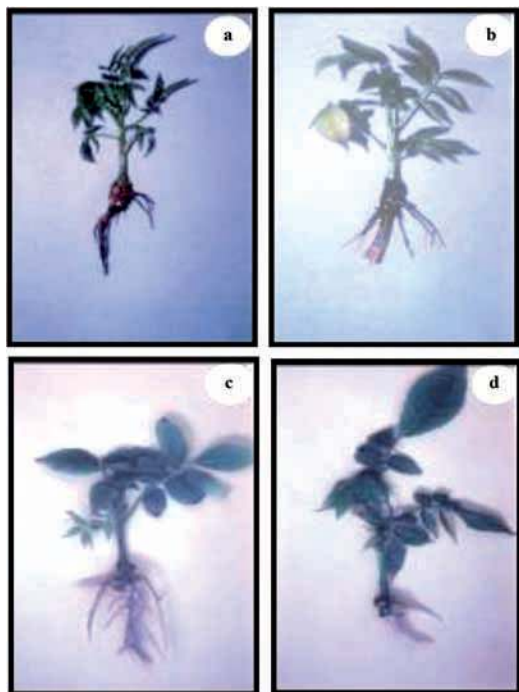


Figure 1. Rooting response in *Juglans regia* L. cv. Serr microshoots. Rooting in T1 and T2 (a and b), T3 and T4 (c and d), respectively.

important difference was that these roots also rapidly produced a high number of secondary roots exceeding the microplants produced with T4 (Figure 1d), characteristic that favors its subsequent acclimatization *ex vitro*.

DISCUSSION

The main problems in the *in vitro* introduction of *J. regia* adult material were related to the permanent appearance of bacterial contamination of endogenous origin called latent contamination (McGranahan *et al.*, 1988), and the exudation of phytotoxic compounds of phenolic origin (Leslie and McGranahan, 1992). However, the use of reinvigorated material through consecutive pruning or macrografting (Claudot *et al.*, 1992; Leslie and McGranahan, 1992) allowed a decrease in the incidence of this problem on the subsequent culture and development of *in vitro* explants (McGranahan *et al.*, 1988), originating synthesis processes of plant growth regulators in quantity and quality similar to that obtained in material of embryonic origin (Sánchez-Olate *et al.*, 2002).

Studies related to this topic concur that the apices portion of the microshoots are the most adequate for the rooting phase, whereas multiplication is better with basal segments of microshoots (Ríos *et al.*, 2002; Sánchez-Olate *et al.*, 2002).

By means of auxinic induction, the rooting rates were greater and the quality of the radicular system gave better results with respect to the number of roots and the zone from which it originated in the microshoot. Rooting was observed at 67.5% compared to 50% attained with *A. rhizogenes*, these values being similar to those obtained by Caboni *et al.* (1996) in embryogenic tissue, condition which can indicate that these bacterial strains have the capacity to improve under organogenic potential imposed by ontogeny (Rodríguez *et al.*, 2005). Infection with *A. rhizogenesis* in the base of the microcuttings was able to induce adventitious radicular systems similar to those obtained in other recalcitrant species (Damiano and Monticelli, 1998; Gutiérrez-Pesce *et al.*, 1998; Hoshino and Mil, 1998; Pérez-Molphe and Ochoa-Alejo, 1998). Abundant adventitious roots were induced in the cut zone, observing characteristics of the transformed roots for the purpose of bacteria plasmid (Tepfer, 1984; Petit *et al.*, 1986; Narasu and Giri, 2000), showing radicular systems distinct from the A-477 and A-478 strains, where the latter appears to have a greater induction capacity than the former, resulting in the particular differences of each strain (Vahdati *et al.*, 2002; Kaur *et al.*, 2006). Furthermore, the transformed roots were able to regenerate transgenic plants or clones that are viable, genetically stable (Narasu and Giri, 2000), and phenotypically normal (Sánchez-Olate *et al.*, 1997). This could be indicating the interaction between endogenous auxins and the *A. rhizogenes* effect (Falasca *et al.*, 2000).

In spite of the observed differences in the rhizogenic rate obtained via IBA or *A. rhizogenes*, the use of the bacterial vector can be a powerful tool to reproduce selected cultivars that are advanced in age, or recuperate high-value cultivars in a state of deterioration, given that the results obtained with this inductor were comparable to those obtained with microshoots of embryonic origin (Ripetti *et al.*, 1994; Caboni *et al.*, 1996).

Finally, considering that the commercial importance of *J. regia* generates a high demand for grafted plants with germoplasma quality, and that the results of the grafting programs are significantly less than the rooting percentages achieved in this study, the use of rhizogenic induction agents can mean plant conversion at higher rates than those achieved via traditional grafting.

CONCLUSIONS

It is possible to attain rooting rates of 50% from *Juglans regia* adult material previously rejuvenated through grafting by using auxinic inducers and *Agrobacterium rhizogenes*. The highest percentage of rooting was obtained in auxinic induction treatments.

RESUMEN

Inducción rizogénica en tejido adulto de *Juglans regia* L. cv. Serr mediada por ácido indol butírico y *Agrobacterium rhizogenes*. La introducción *in vitro* de tejido adulto de nogal (*Juglans regia* L.) representa una oportunidad de clonación de genotipos elite, cuya selección ocurre en estados ontogénicos avanzados. Así, con el objeto de desarrollar un protocolo que permita la propagación masiva de genotipos valiosos a partir de material adulto, se compararon dos sistemas de inducción rizogénica de microtallos de nogal provenientes del cuarto subcultivo de un programa de introducción *in vitro* de tejido adulto de nogal, previamente revigorizado mediante injerto tradicional. Se utilizó la inducción rizogénica por ácido indol-3-butírico (AIB) y *Agrobacterium rhizogenes*. El proceso rizogénico se analizó tanto para inducción auxínica (T1: 3 mg L⁻¹ AIB; T2: 5 mg L⁻¹ AIB), como para inducción por *A. rhizogenes* (T3: A-477; T4: A-478), en dos fases. Una primera fase de inducción radicular, con una duración de 3 días en oscuridad; y una segunda fase de 27 días, denominada de manifestación radicular. En todos los tratamientos se evaluó porcentaje de enraizamiento y se caracterizaron los sistemas radiculares inducidos (número, largo, diámetro y zona de inserción de raíces). Los mejores resultados de enraizamiento se obtuvieron en T2; sin embargo, la respuesta obtenida con *A. rhizogenes* no difiere de aquella lograda en T1, por lo que pareciera ser una metodología de creciente interés para la rizogénesis adventicia en esta especie.

Palabras clave: enraizamiento, microtallos, material adulto, *Agrobacterium rhizogenes*.

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LIFE, SEASONAL CYCLES, AND POPULATION FLUCTUATION OF *Hippodamia variegata* (GOEZE) (COLEOPTERA: COCCINELLIDAE), IN THE CENTRAL PLAIN OF LA ARAUCANÍA REGION, CHILE

Ramón Rebolledo¹*, Johnny Sheriff¹, Leonardo Parra¹, and Alfonso Aguilera¹

ABSTRACT

This study was performed on an alfalfa crop located on the central plain of La Araucanía Region, Chile and in the Laboratorio de Entomología Aplicada de la Facultad de Ciencias Agropecuarias y Forestales at the Universidad de La Frontera. Certain aspects of the biology of *Hippodamia variegata* (Goeze) (Coleoptera: Coccinellidae) were determined, more specifically in relation to its life cycle, seasonality, and population fluctuation. It was established that this coccinellid requires 190.32 ± 10.2 degree-days to complete a generation under laboratory conditions. This information along with the field samplings made it possible to calculate that *H. variegata* completes four generations per season in the alfalfa crop (*Medicago sativa* L.).

Key words: life cycle, seasonal cycle, population fluctuation, *Hippodamia variegata*, *Medicago sativa*.

INTRODUCTION

Hippodamia variegata (Goeze) is an active aphid predator used in the biological control of plant lice in cereals and oil plants in diverse countries (Linskii, 1984; Zúñiga, 1985; Zúñiga *et al.*, 1986; Obrycki and Orr, 1990; Shing and Shing, 1994; El-Hag and Zaitoon, 1996; Obrycki, 1998; González, 2006). Its origin is Palearctic, with a cosmopolitan distribution (Krafsur *et al.*, 1996; Franzmann, 2002), and is found in Asia (Kim *et al.*, 1968; Butani, 1972; Hameed *et al.*, 1977; Wu, 1986), Africa (Badawy, 1969; Haile and Megenasa, 1987; Aalbersberg *et al.*, 1988; Saharaoui and Gourreau, 1998), and Europe (Pruszyński and Lipa, 1971; Natskova, 1973; Radwan and Lovei, 1982; García and Ribeiro, 1983; Plaza, 1987; Ferran *et al.*, 1989; Nicoli *et al.*, 1995; Pekín, 1996; Burgio *et al.*, 2006). It was first introduced in Chile in 1967 as a result of the manifestation of the pale green louse of *Metopolophium dirhodum* (Walk.) gramineae and the dark ear louse of *Sitobion avenae* (Fabricius) (Rojas, 1980 a; 1980b). *H. variegata* is found in Chile from the Arica and Parinacota Region to the Los Lagos Region (González, 2006). According to Aguilera *et al.* (2005, 2006) and Rebolledo *et al.* (2007), its occurrence is notable in La Araucanía Region and is very abundant. Grigorov

(1977), Honek (1985) and Rebolledo *et al.* (2007) state that *H. variegata* prefers herbaceous plants. Nevertheless, Rebolledo *et al.* (2007) point out that it is possible to find this species in shrubby and arboreous plants.

Hagen (1962) affirms that coccinellids determine their conduct through four fundamental actions: voltinism, dormancy or diapause, migrations, and formation of aggregates (Hagen, 1962; Hodek, 1967). Voltinism (the number of generations per year) varies according to latitude. Hagen (1962) recognizes four types of voltinism: I = one generation; II = two generations; III or IV = three or more generations; IA = one generation whose adults migrate to hibernate.

Diapause is intimately related to voltinism given that the latter is a consequence of the former (Nieto y Mier, 1985). Hagen (1962) states that there are three types of dormancy depending on the season: (1) hibernation (Types with voltinism I, II, and III), (2) estivation and hibernation (Type IIA), (3) estivo-hibernation (Type IA). With regards to the formation of aggregates, this same author points out that this is perhaps the most fascinating phenomenon of the coccinellids. The majority of them have an instinctive tendency to hibernate socially, just as it occurs with established tribes such as Hippodamini and Anisocictini. *H. variegata* prefer an aggregation site on mountain tops or close to these (Khan *et al.*, 2007). With respect to the migration phenomenon, Hagen (1962) indicates that the long migratory flights are related to the search for dormancy sites, and that these are associated at the same time with the formation of aggregates. This coccinellid has been studied in the country as regards

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distribution (Arias, 2000) and predatory activity (Grez and Prado, 2000; Grez and Villagrán, 2000). However, there are no data about voltinism and population fluctuation. In La Araucanía Region, *H. variegata* is an abundant species and especially in alfalfa (*Medicago sativa* L.) (Rebolledo *et al.*, 2007).

In order to complement the previous studies and increase knowledge about its behavior in La Araucanía, it was proposed to determine its life and seasonal (voltinism) cycles, fluctuation, relative abundance, and its possible natural entomophagous enemies.

MATERIALS AND METHODS

Field work

Two seasons (1999-2000 and 2000-2001) were required to determine fluctuation and population density of *H. variegata* (in an alfalfa field located in the former Estación Experimental Maipo belonging to the Facultad de Ciencias Agropecuarias y Forestales of the Universidad de La Frontera, located in the urban radius of the city of Temuco (38°44' S, 72°35' W, 100 m.a.s.l.). Breeding of *H. variegata* adults was done with the alfalfa greenbug *Acyrtosiphon pisum* (Harris).

H. variegata adults were periodically collected with entomological nets during October and March in an alfalfa field. Adults were collected in December and were placed in plastic containers measuring 6.5 cm in height, 5.0 cm diameter, and covered with tulle.

To determine the relative abundance and population fluctuation between October 1999 and March 2001, the alfalfa field was visited 46 times. The sampling was carried out periodically every 10 ± 1 day during the spring, summer, and autumn months, and every 20 ± 1 day during the winter months. Each sample consisted in passing 20 times with a standard 30 cm diameter entomological net over the foliage in a 180° range, at a regular pace following the methodology proposed by Metcalf and Luckman (1990) and Apablaza and Stevenson (1995). The sampled area was divided into four 1 ha⁻¹ quadrants, numbered clockwise to facilitate sampling. Three samples were taken in each quadrant by leaving a minimum distance of 25 m between each replicate, checking, and registering the collected material. To determine the number of individuals per m², considering that the width was the amplitude that the net covered in a 180° horizontal movement and the length as the distance covered in the sample.

Laboratory work

H. variegata adults collected in the field were moved to a germination chamber (Archiclíma, Temuco, Chile) with controlled humidity conditions ($70 \pm 8\%$), temperature (21 ± 2 °C), and photoperiod (16:8 light:darkness) to observe

their behavior. Following copulation, males and females were separated. The confined adults were controlled daily to register and eliminate the parasitoids from the breeding.

The laboratory study of the *H. variegata* life cycle was initiated by obtaining eggs from the adults collected in December. These were deposited on Petri dishes and were incubated in the germination chamber in the above-mentioned conditions. When the eggs hatched, 80 larvae were separated, individually placed on numbered plastic dishes, and named initial breeding or group A. From this initial breeding, 57 adults were obtained from which 28 couples were formed at the beginning of January and separated into two groups of 14. A subgroup called AI was made up of 14 females that were permanently maintained with a male, and the second subgroup (AII) was formed by the remaining couples, but maintained with the male only during 48 h. Ten eggs were taken from each female which were bred in isolation in order to determine the influence of the male in the oviposition.

To measure the duration of the life cycle during the month of January, 48 individualized larvae were used as group B and obtained from the 10 isolated eggs of each couple. A new group of 40 larvae were chosen randomly from a single emergence date to determine the growth of each larval stage. The measurements were taken and registered every 24 h. Seven larvae from each larval stage were placed in glass containers with 75% alcohol to subsequently measure the length and width of each one with the help of graph paper. The widest sector of the thorax was used to measure the width, and the length was considered from the top of the head to where the abdomen ends. Furthermore, the width and length of seven pupas, also chosen randomly from groups A and B, were measured.

From the individuals used previously, 22 adults were taken to determine their longevity (50% males and 50% females), were fed daily with *A. pisum* and observed until the moment of their natural death.

The summation of degree-days required for a generation was established to determine the seasonal cycle of *H. variegata*. The formula proposed by Dinelli (1999) was used to calculate this summation. This value was contrasted with the results obtained during the months of field study allowing to determine the degree-days required for the activation of the adults following their hibernation, and the number of generations that theoretically occurred in the study zone.

The reason to use two groups (A and B) was the fact that these coccinellids were collected in different periods, and hence separate statistical analyses were done. An experimental randomized complete block design was used where each individual corresponded to one replicate.

The data of the specimens that completed the life cycle (groups A and B) were compared with variance analysis. Then these results were analyzed using appropriate tests to compare the means of two independent samples, whether parametric (t-Student) or nonparametric (U - Mann-Whitney) (Visauta, 1997; 1998).

RESULTS AND DISCUSSION

Population fluctuation of *H. variegata*

The first adults were examined during October and their number varied between seasons for what seemed to be a clear dependence on the particular yearly environmental conditions. They were completely absent during the winter which accounts for their natural behavior to enter diapause or hibernation. The fluctuation during the period under study showed two annual maxima, one on 10 December 1999 (15 specimens) and the second on 6 February 2001 (13 specimens) (Figure 1).

A noticeable increase in population levels occurred at the beginning of autumn, possibly related with the generation that spent the winter in diapause, results which coincided with those obtained by Apablaza and Stevenson (1995) who pointed out that the annual population maxima for coccinellids in alfalfa in the Metropolitana Region took place at the end of March.

Relative abundance of *H. variegata*

The relative abundance of this insect was 0.17 ± 0.18 adult specimens m^{-2} with a variation of 0 to 0.68 individuals m^{-2} . The larvae showed a mean relative abundance of 0.16 ± 0.25 specimens m^{-2} fluctuating between 0 and 1 individuals m^{-2} .

Volitinism of *H. variegata*

H. variegata required 190.32 ± 10.2 degree-days to complete a generation considering $10\text{ }^{\circ}\text{C}$ as the threshold temperature, the one recommended for the majority of coccinellids. During the first sampling season (spring 1999-summer 2000), it was difficult to estimate the number of generations due to a generational overlap since larvae and adults were found in the crop at the beginning of the measurements. Taking into account only the degree-days accumulated during October and November, it was estimated that the necessary degrees would be on 28 November to complete a first generation. The second generation would be obtained at the end of December, a third on the first days of February, and a fourth and final generation during the first week of March, going through the winter in diapause as an adult.

In the second season (spring 2000-summer 2001), the exact date was registered when adults first appeared coming out of their diapause. It was estimated that soon after the appearance of posthibernational adults, at the end of September, the first generation in the crop was completed around 10 December, obtaining the second generation during the second half of January, and the third generation at the end of the first half of February (Figure 2).

These results differed from those informed by Kontodimas and Stathas (2005) who indicate that a study carried out in Greece using *H. variegata* as a food source for *D. crataegi*, completed a total of seven generations between April and November.

H. variegata completed four generations during this study under the environmental conditions of the central plain of La Araucanía Region, the first of them with a longer duration with two to two and one half months, whereas

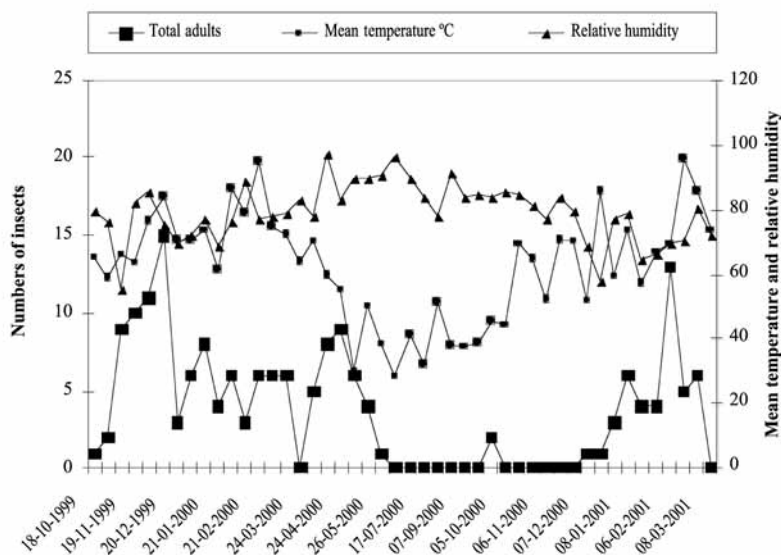


Figure 1. Population fluctuation of *Hippodamia variegata* adults in the central plain of La Araucanía Region.

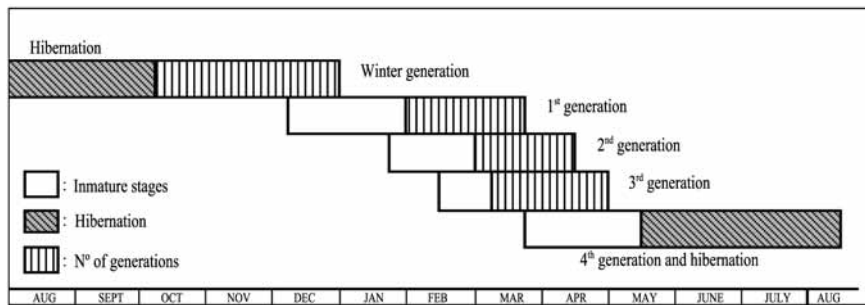


Figure 2. Diagram of *Hippodamia variegata* voltinism in the central plain of La Araucanía Region (2000-2001).

the rest only took one month that is, they showed a type III voltinism in accordance with that proposed by Hagen (1962). The fact that there was a high overlap between one generation and another is emphasized, fact essentially due to the extensive duration of the adult stage in comparison to the rest of the stages of the life cycle. The polyvoltinism observed concurred with Hagen (1962) who pointed out that the environmental conditions are what determine the number of possible generations for coccinellids.

Life cycle of *H. variegata*

H. variegata showed a life cycle of 17.3 ± 0.93 days varying in a range of 16 to 21 days. The pupa stage showed the longest duration with 32% of the life cycle total time, followed by the egg stage and fourth larval stage with 17% each. There is a discrepancy between this result and the one reported by Badawy (1969) who indicates a mean duration of 10.7 days for the life cycle. Breeding temperature would be a fundamental factor in the rate of preimaginal development according to Hagen (1962), Mitchels and Bateman (1986) who mention a duration of 15.1 days at 25 °C for the *H. variegata* life cycle, whereas the life cycle decreased to 7.8 days at 30 °C. The difference between this study and those stated by Hagen (1962) and Badawy (1969) can be attributed to the use of other breeding temperatures.

Oviposition, hatching, and incubation period of *H. variegata* eggs

Mating occurred between 2 and 5 days of life, registering the first ovipositions two days later, which is in accordance with that pointed out by Hodek (1967) and Badawy (1969). The oviposition and hatching of the *H. variegata* eggs are shown in Table 1. With respect to incubation time, it was three days for both groups.

The mean monthly oviposition of *H. variegata* was 223 ± 103.9 eggs and its frequency reached 4.37 ± 5.82 days, a contrast with that reported by Kontodimas and Stathas (2005) who obtained a mean of 956.6 eggs in one breeding of *H. variegata* carried out in Greece. However, that study

used *Dysaphis crataegi* (Kaltenbach) as a food source at a temperature of 25 °C, while the temperature in the present research was 21 °C and the food was *Acyrtosiphon pisum* (Harris). The mean of eggs per day of oviposition in the AI subgroup reached 17.58 ± 6.73 , whereas the groups of eggs attained a mean of 1.59 ± 0.41 per day of oviposition. At the same time, subgroup AII showed values similar to those in the previous subgroup with a mean oviposition of 18.73 ± 6.33 eggs per day of oviposition, a mean of 1.61 ± 0.27 groups per day of oviposition. Differences were not significant between the two subgroups.

Duration of *H. variegata* larval stages

In group A, made up of larvae emerged from the AI and AII subgroups, the mean duration of their development was 7.98 ± 0.73 , days with a range that fluctuated between 7 and 10 days, with the first and fourth stages having the longest duration with 26 and 32%, respectively, considering the total development of this stage of the insect, and coinciding with that reported by Badawy (1969). However, it must be pointed out that this author used the aphid *Aphis gossypii* Glover as food for the *H. variegata* larvae.

The larval stage in group B developed in 10.67 ± 1.62 days with a range that varied between 9 and 15 days, coinciding with that found by Mitchels and Flanders (1992), El-Hag and Zaitoon (1996) who determined a mean duration of the larval stage of 11 days using *Diuraphis noxia* (Mordvilko) and *Brevicoryne brassicae* (L.) + *Rhopalosiphum padi* (L.) as food, respectively. This would indicate that the aphid species used did not determine a difference in the duration of the larval stage. The difference in the duration of the larval period between groups A and B would be due to the capture period of the adults (December for group A and January for group B) in which breeding was initiated (Figure 3). The results of the growth of the randomly chosen larval stages are shown in Table 2.

Duration of *H. variegata* pupa stage

The duration of the pupa stage showed significant differences depending on its origin as group A or B.

Table 1. Oviposition and egg hatching percentage of *Hippodamia variegata*.

Subgroups	Eggs day ⁻¹	Hatching mean (%)	Groups of eggs day ⁻¹	variation (%)
AI ¹	17.58 ± 6.73	87.81 ± 5.76	1.59 ± 0.41	80-100
AII ²	18.73 ± 6.33	81.96 ± 7.27	1.61 ± 0.27	72-100

¹Permanent couple. ²Couple with male only 48 h.

Table 2. Larval instar size of *Hippodamia variegata*.

State	Mean length (mm)	Range	Mean width (mm)	Range
1	1.64 ± 0.56	1-1.15	0.5	0
2	2.28 ± 0.27	2-2.25	1	0
3	4.07 ± 0.45	3.5-4.5	2.28 ± 0.27	2-2.25
4	6.57 ± 0.44	6-7	2	2.25

Group A had a mean duration of 6.27 ± 0.45 days with a range of 6 to 7 days, while group B reached a mean of 4.28 ± 0.61 days and a variation between 3 and 5 days. Badawy (1969) points out a smaller mean duration value of 2.61 days than the one found in this study, difference based on a higher breeding temperature. The mean size of the randomly chosen pupas was a length of 4.57 ± 0.45 mm varying between 4 and 5 mm, and an observed width of 2.57 ± 0.45 with a range of 2 to 3 mm.

Longevity of *H. variegata*

The longevity mean of the adults in both groups was 53.27 ± 11.93 days. The females showed longevity of 55.09 ± 10.85 days and the males 51.45 ± 13.03 days. However, these numbers did not show any significant differences. The mean longevity attained in this study was less than the one cited by El-Hag and Zaitoon (1996) who

fed *H. variegata* with *B. brassicae* and *R. padi* at 25 ± 2 °C, and observed a mean adult longevity of 70 days.

Natural enemies of *H. variegata*

The only species of parasitoid found was *Dinocampus coccinellae* (Schrank) (Hymenoptera: Braconidae) which affected 30% of the adults collected in the field according to the breeding mortality register of adults collected in alfalfa.

CONCLUSIONS

The life cycle of *H. variegata* had a mean duration of 17.3 ± 0.93 days, with a mean adult longevity of 53.27 ± 10.82 days. Both measurements did not differ statistically when comparing both groups grupos ($P = 0.377$ and $P = 0.485$ for the life cycle and longevity, respectively).

It was determined that *H. variegata* required 190.32 ± 10.2 degree-days to complete a generation, signifying that under the existing environmental conditions in the central plain of La Araucanía Region, this coccinellid can complete up to four generations per season. However, based on the population fluctuation and relative abundance records, it is concluded that this insect is not abundant as a natural control agent in the zone under study.

The population fluctuation of coccinellid was markedly seasonal with a complete absence of specimens during the winter and progressive population increases up to an annual maximum at the beginning and middle of the summer, during the first and second year, respectively. The relative abundance of this insect in alfalfa was 0.17 ± 0.18 specimens m⁻² for adults and 0.16 ± 0.25 specimens m⁻² for larvae.

The presence of the hymenoptera parasitoid *Dinocampus coccinellae* (Schrank) (Hymenoptera: Braconidae) was confirmed in 30% of the adults collected in the alfalfa crop.

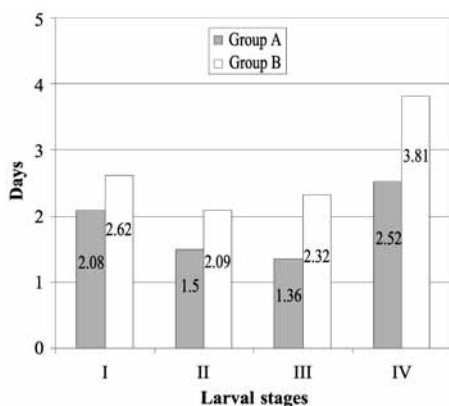


Figure 3. Duration (days) of the *Hippodamia variegata* larval stages under artificial breeding conditions. Group A (December) and Group B (January).

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RESUMEN

Ciclo vital, estacional y fluctuación poblacional de *Hippodamia variegata* (Goeze) (Coleoptera: Coccinellidae), en el llano central de La Araucanía, Chile. El presente estudio fue llevado a cabo en un cultivo de alfalfa ubicado en el llano central de la Región de La Araucanía, Chile, y en el laboratorio de Entomología Aplicada de la Facultad de Ciencias Agropecuarias y Forestales de la Universidad de La Frontera, donde se determinaron aspectos de la biología de *Hippodamia variegata* (Goeze) (Coleoptera: Coccinellidae), específicamente en relación a su ciclo vital, estacional y fluctuación poblacional. Se determinó que en condiciones de laboratorio este coccinélido requiere $190,32 \pm 10,2$ grados días para completar una generación, antecedente que sumado a los muestreos de campo permitió estimar que *H. variegata* completa cuatro generaciones por temporada en el cultivo de alfalfa (*Medicago sativa* L.).

Palabras clave: ciclo vital, ciclo estacional, fluctuación poblacional, *Hippodamia variegata*, *Medicago sativa*.

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LIFE, SEASONAL CYCLES, AND POPULATION FLUCTUATION OF *Hippodamia variegata* (GOEZE) (COLEOPTERA: COCCINELLIDAE), IN THE CENTRAL PLAIN OF LA ARAUCANÍA REGION, CHILE

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ABSTRACT

This study was performed on an alfalfa crop located on the central plain of La Araucanía Region, Chile and in the Laboratorio de Entomología Aplicada de la Facultad de Ciencias Agropecuarias y Forestales at the Universidad de La Frontera. Certain aspects of the biology of *Hippodamia variegata* (Goeze) (Coleoptera: Coccinellidae) were determined, more specifically in relation to its life cycle, seasonality, and population fluctuation. It was established that this coccinellid requires 190.32 ± 10.2 degree-days to complete a generation under laboratory conditions. This information along with the field samplings made it possible to calculate that *H. variegata* completes four generations per season in the alfalfa crop (*Medicago sativa* L.).

Key words: life cycle, seasonal cycle, population fluctuation, *Hippodamia variegata*, *Medicago sativa*.

INTRODUCTION

Hippodamia variegata (Goeze) is an active aphid predator used in the biological control of plant lice in cereals and oil plants in diverse countries (Linskii, 1984; Zúñiga, 1985; Zúñiga *et al.*, 1986; Obrycki and Orr, 1990; Shing and Shing, 1994; El-Hag and Zaitoon, 1996; Obrycki, 1998; González, 2006). Its origin is Palearctic, with a cosmopolitan distribution (Krafsur *et al.*, 1996; Franzmann, 2002), and is found in Asia (Kim *et al.*, 1968; Butani, 1972; Hameed *et al.*, 1977; Wu, 1986), Africa (Badawy, 1969; Haile and Megenasa, 1987; Aalbersberg *et al.*, 1988; Saharaoui and Gourreau, 1998), and Europe (Pruszyński and Lipa, 1971; Natskova, 1973; Radwan and Lovei, 1982; García and Ribeiro, 1983; Plaza, 1987; Ferran *et al.*, 1989; Nicoli *et al.*, 1995; Pekín, 1996; Burgio *et al.*, 2006). It was first introduced in Chile in 1967 as a result of the manifestation of the pale green louse of *Metopolophium dirhodum* (Walk.) gramineae and the dark ear louse of *Sitobion avenae* (Fabricius) (Rojas, 1980 a; 1980b). *H. variegata* is found in Chile from the Arica and Parinacota Region to the Los Lagos Region (González, 2006). According to Aguilera *et al.* (2005, 2006) and Rebolledo *et al.* (2007), its occurrence is notable in La Araucanía Region and is very abundant. Grigorov

(1977), Honek (1985) and Rebolledo *et al.* (2007) state that *H. variegata* prefers herbaceous plants. Nevertheless, Rebolledo *et al.* (2007) point out that it is possible to find this species in shrubby and arboreous plants.

Hagen (1962) affirms that coccinellids determine their conduct through four fundamental actions: voltinism, dormancy or diapause, migrations, and formation of aggregates (Hagen, 1962; Hodek, 1967). Voltinism (the number of generations per year) varies according to latitude. Hagen (1962) recognizes four types of voltinism: I = one generation; II = two generations; III or IV = three or more generations; IA = one generation whose adults migrate to hibernate.

Diapause is intimately related to voltinism given that the latter is a consequence of the former (Nieto y Mier, 1985). Hagen (1962) states that there are three types of dormancy depending on the season: (1) hibernation (Types with voltinism I, II, and III), (2) estivation and hibernation (Type IIA), (3) estivo-hibernation (Type IA). With regards to the formation of aggregates, this same author points out that this is perhaps the most fascinating phenomenon of the coccinellids. The majority of them have an instinctive tendency to hibernate socially, just as it occurs with established tribes such as Hippodamini and Anisocictini. *H. variegata* prefer an aggregation site on mountain tops or close to these (Khan *et al.*, 2007). With respect to the migration phenomenon, Hagen (1962) indicates that the long migratory flights are related to the search for dormancy sites, and that these are associated at the same time with the formation of aggregates. This coccinellid has been studied in the country as regards

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distribution (Arias, 2000) and predatory activity (Grez and Prado, 2000; Grez and Villagrán, 2000). However, there are no data about voltinism and population fluctuation. In La Araucanía Region, *H. variegata* is an abundant species and especially in alfalfa (*Medicago sativa* L.) (Rebolledo *et al.*, 2007).

In order to complement the previous studies and increase knowledge about its behavior in La Araucanía, it was proposed to determine its life and seasonal (voltinism) cycles, fluctuation, relative abundance, and its possible natural entomophagous enemies.

MATERIALS AND METHODS

Field work

Two seasons (1999-2000 and 2000-2001) were required to determine fluctuation and population density of *H. variegata* (in an alfalfa field located in the former Estación Experimental Maipo belonging to the Facultad de Ciencias Agropecuarias y Forestales of the Universidad de La Frontera, located in the urban radius of the city of Temuco (38°44' S, 72°35' W, 100 m.a.s.l.). Breeding of *H. variegata* adults was done with the alfalfa greenbug *Acyrtosiphon pisum* (Harris).

H. variegata adults were periodically collected with entomological nets during October and March in an alfalfa field. Adults were collected in December and were placed in plastic containers measuring 6.5 cm in height, 5.0 cm diameter, and covered with tulle.

To determine the relative abundance and population fluctuation between October 1999 and March 2001, the alfalfa field was visited 46 times. The sampling was carried out periodically every 10 ± 1 day during the spring, summer, and autumn months, and every 20 ± 1 day during the winter months. Each sample consisted in passing 20 times with a standard 30 cm diameter entomological net over the foliage in a 180° range, at a regular pace following the methodology proposed by Metcalf and Luckman (1990) and Apablaza and Stevenson (1995). The sampled area was divided into four 1 ha⁻¹ quadrants, numbered clockwise to facilitate sampling. Three samples were taken in each quadrant by leaving a minimum distance of 25 m between each replicate, checking, and registering the collected material. To determine the number of individuals per m², considering that the width was the amplitude that the net covered in a 180° horizontal movement and the length as the distance covered in the sample.

Laboratory work

H. variegata adults collected in the field were moved to a germination chamber (Archiclíma, Temuco, Chile) with controlled humidity conditions ($70 \pm 8\%$), temperature (21 ± 2 °C), and photoperiod (16:8 light:darkness) to observe

their behavior. Following copulation, males and females were separated. The confined adults were controlled daily to register and eliminate the parasitoids from the breeding.

The laboratory study of the *H. variegata* life cycle was initiated by obtaining eggs from the adults collected in December. These were deposited on Petri dishes and were incubated in the germination chamber in the above-mentioned conditions. When the eggs hatched, 80 larvae were separated, individually placed on numbered plastic dishes, and named initial breeding or group A. From this initial breeding, 57 adults were obtained from which 28 couples were formed at the beginning of January and separated into two groups of 14. A subgroup called AI was made up of 14 females that were permanently maintained with a male, and the second subgroup (AII) was formed by the remaining couples, but maintained with the male only during 48 h. Ten eggs were taken from each female which were bred in isolation in order to determine the influence of the male in the oviposition.

To measure the duration of the life cycle during the month of January, 48 individualized larvae were used as group B and obtained from the 10 isolated eggs of each couple. A new group of 40 larvae were chosen randomly from a single emergence date to determine the growth of each larval stage. The measurements were taken and registered every 24 h. Seven larvae from each larval stage were placed in glass containers with 75% alcohol to subsequently measure the length and width of each one with the help of graph paper. The widest sector of the thorax was used to measure the width, and the length was considered from the top of the head to where the abdomen ends. Furthermore, the width and length of seven pupas, also chosen randomly from groups A and B, were measured.

From the individuals used previously, 22 adults were taken to determine their longevity (50% males and 50% females), were fed daily with *A. pisum* and observed until the moment of their natural death.

The summation of degree-days required for a generation was established to determine the seasonal cycle of *H. variegata*. The formula proposed by Dinelli (1999) was used to calculate this summation. This value was contrasted with the results obtained during the months of field study allowing to determine the degree-days required for the activation of the adults following their hibernation, and the number of generations that theoretically occurred in the study zone.

The reason to use two groups (A and B) was the fact that these coccinellids were collected in different periods, and hence separate statistical analyses were done. An experimental randomized complete block design was used where each individual corresponded to one replicate.

The data of the specimens that completed the life cycle (groups A and B) were compared with variance analysis. Then these results were analyzed using appropriate tests to compare the means of two independent samples, whether parametric (t-Student) or nonparametric (U - Mann-Whitney) (Visauta, 1997; 1998).

RESULTS AND DISCUSSION

Population fluctuation of *H. variegata*

The first adults were examined during October and their number varied between seasons for what seemed to be a clear dependence on the particular yearly environmental conditions. They were completely absent during the winter which accounts for their natural behavior to enter diapause or hibernation. The fluctuation during the period under study showed two annual maxima, one on 10 December 1999 (15 specimens) and the second on 6 February 2001 (13 specimens) (Figure 1).

A noticeable increase in population levels occurred at the beginning of autumn, possibly related with the generation that spent the winter in diapause, results which coincided with those obtained by Apablaza and Stevenson (1995) who pointed out that the annual population maxima for coccinellids in alfalfa in the Metropolitana Region took place at the end of March.

Relative abundance of *H. variegata*

The relative abundance of this insect was 0.17 ± 0.18 adult specimens m^{-2} with a variation of 0 to 0.68 individuals m^{-2} . The larvae showed a mean relative abundance of 0.16 ± 0.25 specimens m^{-2} fluctuating between 0 and 1 individuals m^{-2} .

Volitinism of *H. variegata*

H. variegata required 190.32 ± 10.2 degree-days to complete a generation considering $10\text{ }^{\circ}\text{C}$ as the threshold temperature, the one recommended for the majority of coccinellids. During the first sampling season (spring 1999-summer 2000), it was difficult to estimate the number of generations due to a generational overlap since larvae and adults were found in the crop at the beginning of the measurements. Taking into account only the degree-days accumulated during October and November, it was estimated that the necessary degrees would be on 28 November to complete a first generation. The second generation would be obtained at the end of December, a third on the first days of February, and a fourth and final generation during the first week of March, going through the winter in diapause as an adult.

In the second season (spring 2000-summer 2001), the exact date was registered when adults first appeared coming out of their diapause. It was estimated that soon after the appearance of posthibernational adults, at the end of September, the first generation in the crop was completed around 10 December, obtaining the second generation during the second half of January, and the third generation at the end of the first half of February (Figure 2).

These results differed from those informed by Kontodimas and Stathas (2005) who indicate that a study carried out in Greece using *H. variegata* as a food source for *D. crataegi*, completed a total of seven generations between April and November.

H. variegata completed four generations during this study under the environmental conditions of the central plain of La Araucanía Region, the first of them with a longer duration with two to two and one half months, whereas

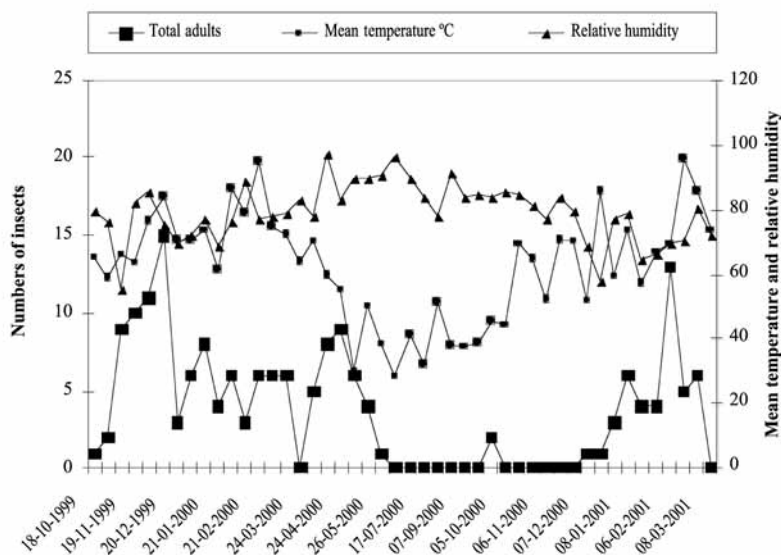


Figure 1. Population fluctuation of *Hippodamia variegata* adults in the central plain of La Araucanía Region.

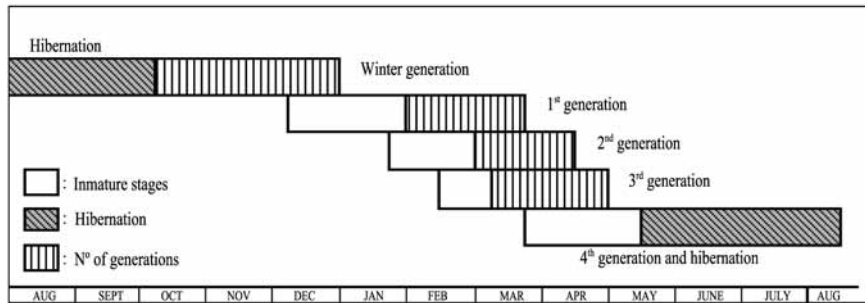


Figure 2. Diagram of *Hippodamia variegata* voltinism in the central plain of La Araucanía Region (2000-2001).

the rest only took one month that is, they showed a type III voltinism in accordance with that proposed by Hagen (1962). The fact that there was a high overlap between one generation and another is emphasized, fact essentially due to the extensive duration of the adult stage in comparison to the rest of the stages of the life cycle. The polyvoltinism observed concurred with Hagen (1962) who pointed out that the environmental conditions are what determine the number of possible generations for coccinellids.

Life cycle of *H. variegata*

H. variegata showed a life cycle of 17.3 ± 0.93 days varying in a range of 16 to 21 days. The pupa stage showed the longest duration with 32% of the life cycle total time, followed by the egg stage and fourth larval stage with 17% each. There is a discrepancy between this result and the one reported by Badawy (1969) who indicates a mean duration of 10.7 days for the life cycle. Breeding temperature would be a fundamental factor in the rate of preimaginal development according to Hagen (1962), Mitchels and Bateman (1986) who mention a duration of 15.1 days at 25 °C for the *H. variegata* life cycle, whereas the life cycle decreased to 7.8 days at 30 °C. The difference between this study and those stated by Hagen (1962) and Badawy (1969) can be attributed to the use of other breeding temperatures.

Oviposition, hatching, and incubation period of *H. variegata* eggs

Mating occurred between 2 and 5 days of life, registering the first ovipositions two days later, which is in accordance with that pointed out by Hodek (1967) and Badawy (1969). The oviposition and hatching of the *H. variegata* eggs are shown in Table 1. With respect to incubation time, it was three days for both groups.

The mean monthly oviposition of *H. variegata* was 223 ± 103.9 eggs and its frequency reached 4.37 ± 5.82 days, a contrast with that reported by Kontodimas and Stathas (2005) who obtained a mean of 956.6 eggs in one breeding of *H. variegata* carried out in Greece. However, that study

used *Dysaphis crataegi* (Kaltenbach) as a food source at a temperature of 25 °C, while the temperature in the present research was 21 °C and the food was *Acyrtosiphon pisum* (Harris). The mean of eggs per day of oviposition in the AI subgroup reached 17.58 ± 6.73 , whereas the groups of eggs attained a mean of 1.59 ± 0.41 per day of oviposition. At the same time, subgroup AII showed values similar to those in the previous subgroup with a mean oviposition of 18.73 ± 6.33 eggs per day of oviposition, a mean of 1.61 ± 0.27 groups per day of oviposition. Differences were not significant between the two subgroups.

Duration of *H. variegata* larval stages

In group A, made up of larvae emerged from the AI and AII subgroups, the mean duration of their development was 7.98 ± 0.73 , days with a range that fluctuated between 7 and 10 days, with the first and fourth stages having the longest duration with 26 and 32%, respectively, considering the total development of this stage of the insect, and coinciding with that reported by Badawy (1969). However, it must be pointed out that this author used the aphid *Aphis gossypii* Glover as food for the *H. variegata* larvae.

The larval stage in group B developed in 10.67 ± 1.62 days with a range that varied between 9 and 15 days, coinciding with that found by Mitchels and Flanders (1992), El-Hag and Zaitoon (1996) who determined a mean duration of the larval stage of 11 days using *Diuraphis noxia* (Mordvilko) and *Brevicoryne brassicae* (L.) + *Rhopalosiphum padi* (L.) as food, respectively. This would indicate that the aphid species used did not determine a difference in the duration of the larval stage. The difference in the duration of the larval period between groups A and B would be due to the capture period of the adults (December for group A and January for group B) in which breeding was initiated (Figure 3). The results of the growth of the randomly chosen larval stages are shown in Table 2.

Duration of *H. variegata* pupa stage

The duration of the pupa stage showed significant differences depending on its origin as group A or B.

Table 1. Oviposition and egg hatching percentage of *Hippodamia variegata*.

Subgroups	Eggs day ⁻¹	Hatching mean (%)	Groups of eggs day ⁻¹	variation (%)
AI ¹	17.58 ± 6.73	87.81 ± 5.76	1.59 ± 0.41	80-100
AII ²	18.73 ± 6.33	81.96 ± 7.27	1.61 ± 0.27	72-100

¹Permanent couple. ²Couple with male only 48 h.

Table 2. Larval instar size of *Hippodamia variegata*.

State	Mean length (mm)	Range	Mean width (mm)	Range
1	1.64 ± 0.56	1-1.15	0.5	0
2	2.28 ± 0.27	2-2.25	1	0
3	4.07 ± 0.45	3.5-4.5	2.28 ± 0.27	2-2.25
4	6.57 ± 0.44	6-7	2	2.25

Group A had a mean duration of 6.27 ± 0.45 days with a range of 6 to 7 days, while group B reached a mean of 4.28 ± 0.61 days and a variation between 3 and 5 days. Badawy (1969) points out a smaller mean duration value of 2.61 days than the one found in this study, difference based on a higher breeding temperature. The mean size of the randomly chosen pupas was a length of 4.57 ± 0.45 mm varying between 4 and 5 mm, and an observed width of 2.57 ± 0.45 with a range of 2 to 3 mm.

Longevity of *H. variegata*

The longevity mean of the adults in both groups was 53.27 ± 11.93 days. The females showed longevity of 55.09 ± 10.85 days and the males 51.45 ± 13.03 days. However, these numbers did not show any significant differences. The mean longevity attained in this study was less than the one cited by El-Hag and Zaitoon (1996) who

fed *H. variegata* with *B. brassicae* and *R. padi* at 25 ± 2 °C, and observed a mean adult longevity of 70 days.

Natural enemies of *H. variegata*

The only species of parasitoid found was *Dinocampus coccinellae* (Schrank) (Hymenoptera: Braconidae) which affected 30% of the adults collected in the field according to the breeding mortality register of adults collected in alfalfa.

CONCLUSIONS

The life cycle of *H. variegata* had a mean duration of 17.3 ± 0.93 days, with a mean adult longevity of 53.27 ± 10.82 days. Both measurements did not differ statistically when comparing both groups grupos ($P = 0.377$ and $P = 0.485$ for the life cycle and longevity, respectively).

It was determined that *H. variegata* required 190.32 ± 10.2 degree-days to complete a generation, signifying that under the existing environmental conditions in the central plain of La Araucanía Region, this coccinellid can complete up to four generations per season. However, based on the population fluctuation and relative abundance records, it is concluded that this insect is not abundant as a natural control agent in the zone under study.

The population fluctuation of coccinellid was markedly seasonal with a complete absence of specimens during the winter and progressive population increases up to an annual maximum at the beginning and middle of the summer, during the first and second year, respectively. The relative abundance of this insect in alfalfa was 0.17 ± 0.18 specimens m⁻² for adults and 0.16 ± 0.25 specimens m⁻² for larvae.

The presence of the hymenoptera parasitoid *Dinocampus coccinellae* (Schrank) (Hymenoptera: Braconidae) was confirmed in 30% of the adults collected in the alfalfa crop.

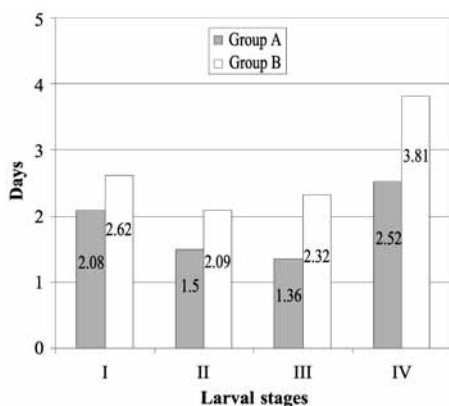


Figure 3. Duration (days) of the *Hippodamia variegata* larval stages under artificial breeding conditions. Group A (December) and Group B (January).

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RESUMEN

Ciclo vital, estacional y fluctuación poblacional de *Hippodamia variegata* (Goeze) (Coleoptera: Coccinellidae), en el llano central de La Araucanía, Chile. El presente estudio fue llevado a cabo en un cultivo de alfalfa ubicado en el llano central de la Región de La Araucanía, Chile, y en el laboratorio de Entomología Aplicada de la Facultad de Ciencias Agropecuarias y Forestales de la Universidad de La Frontera, donde se determinaron aspectos de la biología de *Hippodamia variegata* (Goeze) (Coleoptera: Coccinellidae), específicamente en relación a su ciclo vital, estacional y fluctuación poblacional. Se determinó que en condiciones de laboratorio este coccinélido requiere $190,32 \pm 10,2$ grados días para completar una generación, antecedente que sumado a los muestreos de campo permitió estimar que *H. variegata* completa cuatro generaciones por temporada en el cultivo de alfalfa (*Medicago sativa* L.).

Palabras clave: ciclo vital, ciclo estacional, fluctuación poblacional, *Hippodamia variegata*, *Medicago sativa*.

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