



Probiotic potential of indigenous yeasts isolated during the fermentation of table olives from Northeast of Portugal



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ABSTRACT

The aim of this study was to evaluate the probiotic potential of 16 yeast strains previously isolated during the natural fermentation of table olives *Negrinha de Freixo* cultivar, in relation to enzymatic activities; ability to grow at 37 °C; antimicrobial activity; autoaggregation capacity; antioxidant activity and survival in gastrointestinal tract conditions. The highest antioxidant activity was observed for *Saccharomyces cerevisiae*, similar to the reference strain *S. boulardii*. *Candida norvegica* 7A and *Galactomyces reessii* 34A showed antifungal ability to the pathogenic microorganism *Cryptococcus neoformans*. Regarding the autoaggregation capacity, *S. cerevisiae* 15A, *C. tropicalis* 1A and *C. norvegica* 7A showed > 80% after 24 h. *Pichia guilliermondii* 25A and *C. norvegica* 7A were the most resistant to the simulated digestive conditions, similar to the reference strain (*S. boulardii*). Thus, these results suggest that some yeast strains involved in the fermentation of table olives have probiotic potential.

Industrial relevance: In this study, we highlight the probiotic potential of yeast microbiota usually found in green table olives. These yeast strains could be used as culture starters for the development of new functional products.

1. Introduction

Nowadays the search for a healthier lifestyle has led to the demand by consumers for healthy and natural products that could provide health benefits to the consumers. Accordingly, there has been a breakthrough in the development of so-called probiotics products. Probiotics are defined by the World Health Organization (WHO) as live microorganisms which when administered in adequate amounts confer a health benefit on the host (FAO/WHO, 2006). The term probiotic includes a wide variety of microorganisms, including bacteria and yeast (Perricone, Bevilacqua, Corbo, & Sinigaglia, 2014). The microorganisms often used in probiotic preparations are mostly derived from human sources and/or animals, belonging to the genera *Lactobacillus* and *Bifidobacterium* and less frequently *Enterococcus* (Peres, Peres, Hernández-Mendoza, & Malcata, 2012). The use of these bacteria has been confined almost exclusively to the production of fermented dairy products such as yogurts. However, the increasing number of lactose intolerance cases worldwide, together with the adverse effect of cholesterol content in dairy products (Argyri et al., 2013) and the increased market for vegetarian consumers, has led to the need of plant based probiotic strains that could be used without any restrictions (Rodríguez-Gómez, Romero-

Gil, García-García, Garrido-Fernández, & Arroyo-López, 2014).

According to Ruiz-Moyano, Martín, Benito, Nevado, and Córdoba (2008), traditional fermented products can be a good basis for the development of functional foods of the probiotic type. Thus, the table olives obtained through the natural fermentation process by lactic acid bacteria (LAB) and yeast (natural flora) could be a promising probiotic that can be used as a functional starter culture or as a good source of probiotic microorganism (Argyri et al., 2013). Sourabh, Kanwar, and Sharma (2011) referred that species involved in the fermentation process could be used as protective or probiotic cultures as probiotics are relatively safe for human consumption and are designated as GRAS (*Generally Recognized as Safe*).

Nevertheless, in the last years the search for probiotic microorganisms from olives fermentation has focused on LAB (Argyri et al., 2013; Bautista-Gallego et al., 2013; Pessione, Bianco, Mangiapane, Cirrincione, & Pessione, 2015; Rodríguez-Gómez et al., 2014). On the other hand, there are few studies on yeasts isolated from the fermentation of olives (Bautista-Gallego et al., 2011; Bonatsou, Benítez, Rodríguez-Gómez, Panagou, & Arroyo-López, 2015; Silva et al., 2011). However, there is great interest in finding yeast strains with probiotic potential due to the numerous beneficial effects on human health,

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including prevention and treatment of intestinal diseases such as cholera (Vidon, Huchet, & Rambaud, 1986). Other beneficial functions of yeasts are related with the improvement of bioavailability of minerals through the hydrolysis of phytate (Ragon et al., 2008), anti-inflammatory effects (Mumy, Chen, Kelly, & McCormick, 2008) and detoxification of mycotoxins due to surface binding to the yeast cell wall (Moslehi-Jenabian, Pedersen, & Jespersen, 2010). In addition, yeasts have the ability to produce natural antioxidants such as carotenoids, citric acid, ascorbic acid, tocopherols and glutathione that may retard the oxidative degeneration of lipids and improve the health of the host (Arroyo-López et al., 2012). Furthermore, *Saccharomyces cerevisiae* var. *boulardii* has been proven effective in several double-blind studies and preparations of this yeast have been sold as both a preventive and therapeutic agent (Lukaszewicz, 2012).

Thus, the aim of the present study was to evaluate the probiotic potential of 16 yeast strains isolated and identified on the natural fermentation process of table olives *Negrinha de Freixo* cultivar in relation to the enzymatic activity, ability to grow at 37 °C (human body temperature), autoaggregation capacity, and antimicrobial and antioxidant activities, as well as the survival ability of the strains to the gastrointestinal (GI) transit. The results obtained will be important to promote the use of yeasts with probiotic potential in mixed starter cultures for table olives fermentation and other applications.

2. Material and methods

2.1. Materials

Sodium chloride, potassium dihydrogen phosphate, calcium chloride, potassium chloride, hydrogen peroxide, methanol, hydrochloric acid, disodium hydrogen phosphate, glucose, casein, pepsin, pancreatin and lysozyme were purchased from Panreac (Barcelona, Spain). Potato Dextrose broth, tryptone, yeast extract, agar and Mueller Hinton Agar (MHA) were obtained from Himedia (Kelton, USA). Tween 80, erythromycin, penicillin G and amphotericin B were purchased from Liofilchem (Roseto degli Abruzzi, Italy). Xylan from beech wood and the radical 2,2-diphenyl-1-picryl-hydrazyl (DPPH) were obtained from Sigma Aldrich (St. Louis, USA). Arbutin was purchased from ChemCruz (Huissen, The Netherlands). Casein, phenol red and Brain Heart Infusion broth (BHI) were obtained from Merck (New Jersey, USA).

2.2. Yeast strains

A total of 16 yeast strains usually found in green table olives *Negrinha de Freixo* cv. were used in the present study, namely: *Candida tropicalis* 1A and 3A; *Debaryomyces hansenii* 8A; *Galactomyces reessii* 33A and 34A; *Pichia guilliermondii* 25A; *Pichia manshurica* 2A; *Pichia membranifaciens* 29A and 3B; *Rhodotorula glutinis* 27A; *Rhodotorula graminis* 20A; *Saccharomyces cerevisiae* 15A and 15B; *Candida boidinii* 32A and 37A; and *Candida norvegica* 7A. All of them were previously isolated and identified by Pereira, Ramalhosa, Borges, Pereira, and Baptista (2015) from olives *in natura* (before immersion in brine), olive pulp and brine during the spontaneous fermentation of table olives. The identification of the yeast strains was based on the sequencing of the internal transcribed spacer (ITS) region of the nuclear ribosomal DNA. Its sequences were deposited in NCBI (*National Center for Biotechnology Information*) GenBank data library, under accession numbers KM519627 to KM519637. Lyophilized *Saccharomyces boulardii* strain obtained from the drug Floratil® (Merck, New Jersey, USA) was used as a reference strain.

2.3. Growth at 37 °C and *in vitro* simulated gastric and pancreatic digestions

A pre-inoculum of each strain was initially activated for 24 h in

Potato Dextrose Broth (PDB) at 25 °C, for the study of yeasts' growth. At the end of the incubation period, the optical density (OD) of each culture was measured with the spectrophotometer (Cary 50 Scan), and an aliquot was added to 20 mL broth in order to adjust the OD of the suspension to 0.1 at a wavelength (λ) of 600 nm. Then, the cultures were incubated in a thermostated orbital shaker at two different temperatures (25 and 37 °C), with constant stirring (150 rpm), for 8 h. The OD₆₀₀ values were determined at hour intervals, in order to compare the kinetic growth at both temperatures.

In vitro gastric (GD) and pancreatic digestions (PD) were performed using the methodology described by Bonatsou et al. (2015). For the GD step, a synthetic gastric juice was prepared in a buffer solution at pH 2.0, containing NaCl (2.05 g/L), KH₂PO₄ (0.60 g/L), CaCl₂ (0.11 g/L) and KCl (0.37 g/L), being the pH adjusted with 1 M HCl. After medium's sterilization, pepsin (0.0133 g/L) and lysozyme (0.01 g/L) were added to obtain the gastric solution. The assayed yeast cultures were allowed to grow in PDB (24 h at 25 °C), centrifuged (5000g, 10 min, 4 °C) and the pellet was washed twice with the buffer solution. Afterwards, the cells were re-suspended in the gastric solution, adjusted to an OD value equal to 0.5 (λ = 600 nm) and incubated for 2.5 h at 37 °C, in the orbital shaker at 200 rpm to simulate the peristaltic movements.

The PD simulation was carried out using a buffer solution at pH 8.0, adjusted with 1 M HCl, formulated by bile salts (3.0 g/L), pancreatin (0.1 g/L), Na₂HPO₄ (26.9 g/L) and NaCl (8.5 g/L). Harvested cells from the previous GD step were centrifuged and washed twice in a sterile isotonic solution and re-suspended in the simulated pancreatic solution. After shaking the suspension (200 rpm, 3.5 h at 37 °C), the pellet was washed and the cells were re-suspended in a volume of isotonic solution (0.9% NaCl, w/v). The plating on PDA culture medium, at the beginning and end of each digestive simulation, were performed to assess cell viability. For each strain, four independent assays were performed.

2.4. Enzymatic activity

The enzymatic activity of the yeasts was done using qualitative tests. Strains were initially grown on PDA medium and then inoculated in an enzyme screening medium (except for the catalase test). Each test was carried out in three independent assays and the results were qualitatively expressed as: “–” (no activity); “+” (weak activity); “++” (moderate activity); and “+++” (strong activity). For the catalase, results were expressed in terms of presence “P” or absence “0” of this activity.

Catalase activity of the yeasts was performed according to the method described by Whittenbury (1964), by adding 3% (v/v) hydrogen peroxide onto the yeast colonies. Catalase activity was evidenced by the formation of bubbles.

Protease activity was determined by the method described by Strauss, Jolly, Lambrechts, and Rensburg (2001), by spreading yeast colonies onto PDA containing 20 g/L casein. After the incubation period (5 days at 30 °C), a positive result was considered when a clear zone around the colony was observed.

β -Glucosidase activity assays were performed according to the method described by Bautista-Gallego et al. (2011), using arbutin as substrate. β -Glucosidase activity was evidenced by the appearance of a dark brown colour around the colonies.

Lipase activity was performed according to the method described by Samad et al. (1989), with few modifications. The Yeast Extract Agar medium (YEA, tryptone 6.0 g/L, yeast extract 3.0 g/L and agar 15 g/L), supplemented with Tween 80 (1%) and phenol red (0.01%, w/v) at pH 7.3–7.4, was inoculated with different isolates of yeasts and incubated at 37 °C for 48 h. Lipase activity was observed by the colour change of the medium (red to yellow).

Amylase activity was performed using the medium starch agar. After incubation (30 °C for 5 days), the plates were flooded with iodine solution and the appearance of a brown colour around a colony was

considered a positive result.

The xylanase activity was determined according to the method described by Hernández, Martín, Aranda, Pérez-Navado, and Córdoba (2007). The appearance of a clear halo around the colonies was considered a positive result.

2.5. Antioxidant activity (2,2-diphenyl-1-picryl-hydrazyl (DPPH) free radical scavenging assay)

The assessment of the scavenging ability of the 2,2-diphenyl-1-picryl-hydrazyl (DPPH) radical of each strain was performed in three independent assays, according to the method described by Gil-Rodríguez, Carrascosa, and Requena (2015), with few modifications. Strains were allowed to grow at 37 °C for 24 h, being the OD₆₀₀ adjusted to 1.20, corresponding to a cell concentration on the order of 6–7 log CFU/mL and then centrifuged (7200g, 10 min, 4 °C). The resulting pellet was washed twice with a sterile solution 0.9% NaCl (w/v) and then re-suspended in 2 mL of the same solution. Each test sample (1.6 mL) was added to 2 mL of a DPPH methanolic solution (0.2 mM). After shaking, the mix was incubated for 30 min, at room temperature, in darkness. The scavenging capacity of DPPH free radicals was measured by spectrophotometry (Cary 50 Scan) at $\lambda = 517$ nm. An aliquot containing 1.6 mL of 0.9% NaCl (w/v) and 2 mL of DPPH methanolic solution, which was also incubated for 30 min in the dark, was used as blank. The percentage of antioxidant activity (A.A) was calculated as follows:

$$\%A.A. = [1 - (A_{517}(\text{Sample})/A_{517}(\text{Blank}))] \times 100$$

2.6. Autoaggregation capacity

The autoaggregation percentage of strains was performed using the method described by Gil-Rodríguez et al. (2015). Sample (1.2 mL) of yeast culture in PDB broth was harvested by centrifugation (7200g, 5 min, 4 °C) and washed twice with a sterile solution 0.9% NaCl (w/v). The suspension obtained was re-suspended in 1 mL of the same solution (0.9% NaCl, w/v) and transferred to a disposable plastic cuvette. Without shaking the yeast suspension, the absorbance at 600 nm was measured at the beginning (A_0) and 2, 4 and 24 h (A_T) after that. The autoaggregation percentage was calculated as follows:

$$\% \text{Autoaggregation} = [1 - (A_T/A_0)] \times 100$$

For each strain, three independent assays were performed.

2.7. Antimicrobial activity

All strains were tested in triplicate (independent assays) for antimicrobial activity in the presence of two yeasts (*Candida albicans* and *Cryptococcus neoformans*) and three bacteria (*Pseudomonas aeruginosa*, *Escherichia coli* and *Staphylococcus aureus*) by the agar diffusion method. These pathogenic microorganisms belong to the collection of the School of Agriculture of the Polytechnic Institute of Bragança. The cultures of these microorganisms were inoculated into 5 mL of Brain Heart Infusion broth (BHI) and incubated at 25 °C (yeast) and 37 °C (bacteria) for 16–18 h. Subsequently, the suspensions were diluted into sterile saline (0.9% NaCl, w/v) to obtain a suspension with a turbidity corresponding to 0.5 McFarland standard (8–9 log cfu/mL) and seeded using a sterile swab on the surface of the culture medium (Mueller Hinton Agar (MHA)) for bacteria and MHA with 2% glucose (w/v) for yeast. Once dried, 10 μ L of each suspension of the assayed yeasts were inoculated in paper discs (6.0 mm in diameter, Oxoid), previously placed on the surface of the medium. Antibacterial antibiotics, erythromycin (15 μ g) and penicillin G (10 IU), discs and the antifungal amphotericin B (20 μ g) were used as positive controls. BHI broth at pH 6.5 was used as negative control. The plates were incubated at 37 °C (bacteria) and 25 °C (yeast) during 2–5 days. The results were determined from the

appearance of a clear halo around the filter paper discs.

2.8. Statistical analysis

Statistical analysis of the antioxidant activity, autoaggregation and *in vitro* simulated gastric and pancreatic digestions, was carried out by One-Way Analysis of Variance, being the means compared by the Tukey's test, using SPSS v.20 software (SPSS Inc., Chicago, USA). For determining the specific growth rate (μ) for each of the isolates, a linear regression analysis was carried out using Excel of Microsoft Corporation (2013) and using the values from the natural log of the optical density (LN OD) versus time, comprised in the exponential growth phase.

In order to identify the yeast strains with desirable probiotic properties, Principal Component Analysis (PCA) was applied in the program SPSS (v. 20.0). The results of the qualitative tests (enzyme activity) were converted into four qualitative codes (1 - none; 2 - low, 3 - moderate, 4 - strong activity). For the selection of the number of Principal Components or Factors, the Kaiser criterion test was applied and only factors with eigen-values higher than 1.00 were maintained.

3. Results and discussion

3.1. Growth at 37 °C and *in vitro* simulated gastric and pancreatic digestions

The assessment of the ability of yeast growth at 37 °C is an important parameter when it is desired to select promising probiotic microorganisms (Gil-Rodríguez et al., 2015). To exert their beneficial effects, microorganisms need to multiply within the host. In this study, the strains that showed the highest specific growth rate (μ) at 37 °C (data not shown) were *S. cerevisiae* 15B ($\mu = 0.718 \text{ h}^{-1}$), *C. tropicalis* 1A ($\mu = 0.678 \text{ h}^{-1}$) and *P. membranifaciens* 29A ($\mu = 0.676 \text{ h}^{-1}$), similar to the probiotic reference *S. boulardii* ($\mu = 0.584 \text{ h}^{-1}$). These results showed that this temperature was favorable for their growth and, therefore, these strains can easily adapt to the temperature of the host. Similar results were also obtained by Gil-Rodríguez et al. (2015) for *S. cerevisiae* IFI-1145 and *P. membranifaciens* IFI-946, isolated from food and beverages. In contrast, isolates belonging to the genus *Rhodotorula*, *R. graminis* 20A and *R. glutinis* 27A, showed lower growth rates at 37 °C (0.109 h^{-1} and 0.186 h^{-1} , respectively) than at 25 °C, suggesting growing difficulties at human body temperature. Thus, in our study, only 9 strains of the 16 initially tested strains showed good ability to survive at 37 °C, being afterwards subjected to *in vitro* GD and PD simulation assays, together with *S. boulardii* (reference strain).

Table 1 shows the results of the survival percentage of all tested strains, as well as the probiotic yeast, *S. boulardii*, as reference strain. All strains were resistant to GD process, indicating *C. tropicalis* 3A and *P. guilliermondii* 25A had significantly ($p < 0.05$) higher survival rates than the other strains, including the reference strain. The survival rates to GD of *C. tropicalis* 3A and *P. guilliermondii* 25A were equal to 194.76 \pm 26.52% and 179.43 \pm 25.00%, respectively. These results indicate that these microorganisms are not affected by the environmental conditions of the stomach, as their population almost doubled. The remaining strains showed survival rates ranging between 72.65 \pm 40.55% (*C. tropicalis* 1A) and 3.16 \pm 1.71% (*S. cerevisiae* 15A); however, there was high variability for the same sample, resulting in high standard deviations. Therefore, no significant differences between these samples and the reference strain (44.43 \pm 1.93%) were observed. This variability was also stated by Bonatsou et al. (2015) in a study conducted with yeast isolated from Greek natural black table olives, particularly with *P. guilliermondii* (492.05 \pm 319.63%).

After the period of exposure to GD conditions (2.5 h), the strains were then subjected to PD simulation for 3.5 h. All strains survived to the synthetic pancreatic juice. *S. cerevisiae* 15B showed a value (175.98 \pm 47.84%) significantly higher ($p < 0.05$) than other strains, excluding the strain *C. norvegica* 7A (125.75 \pm 47.24%). These strains increased their effective population, suggesting good adaptability to

Table 1
Survival (%) of the yeast strains to the gastric and pancreatic digestions and overall survival (mean \pm standard deviation, n = 4).

Isolates	Species	Gastric digestion	Pancreatic digestion	Overall survival
1A	<i>C. tropicalis</i>	72.65 \pm 40.55 ^b	56.11 \pm 0.55 ^{b,c}	9.05 \pm 6.40 ^b
3A	<i>C. tropicalis</i>	194.76 \pm 26.52 ^a	61.41 \pm 30.59 ^{b,c}	9.04 \pm 5.91 ^b
15B	<i>S. cerevisiae</i>	39.15 \pm 17.46 ^{b,c,d}	175.98 \pm 47.84 ^a	2.80 \pm 0.36 ^b
15A	<i>S. cerevisiae</i>	3.16 \pm 1.71 ^d	12.61 \pm 11.63 ^c	0.13 \pm 0.13 ^b
7A	<i>C. norvegica</i>	49.75 \pm 13.72 ^{b,c}	125.75 \pm 47.24 ^{a,b}	14.52 \pm 3.32 ^{a,b}
8A	<i>D. hansenii</i>	40.71 \pm 2.03 ^{b,c,d}	97.74 \pm 5.58 ^b	7.38 \pm 3.63 ^b
37A	<i>C. boidinii</i>	16.16 \pm 5.62 ^{c,d}	18.17 \pm 7.75 ^c	0.48 \pm 0.35 ^b
25A	<i>P. guilliermondii</i>	179.43 \pm 25.00 ^a	68.91 \pm 22.59 ^{b,c}	29.13 \pm 16.95 ^a
2A	<i>P. manshurica</i>	29.63 \pm 7.46 ^{b,c,d}	11.05 \pm 6.09 ^c	9.53 \pm 2.98 ^{a,b}
Sb	<i>S. boulardii</i>	44.43 \pm 1.93 ^{b,c,d}	99.79 \pm 34.11 ^b	13.33 \pm 8.98 ^{a,b}

Different letters in the same column indicate significant differences by Tukey's test ($p < 0.05$).

these environmental conditions. Silva et al. (2011) also demonstrated similar results for the yeasts isolated from Portuguese brine table olives to bile salts.

Regarding the overall survival of the simulation of both digestive processes (GD and PD), it was observed that the strain *P. guilliermondii* 25A showed the greatest resistance to the entire digestive process, with a value of 29.13 \pm 16.95%. This value was statistically similar to that of *S. boulardii* and the strains *C. norvegica* 7A (14.52 \pm 3.32%) and *P. manshurica* 2A (9.53 \pm 2.98%). *C. boidinii* 37A and *S. cerevisiae* 15A showed the lowest resistance, with rates of 0.48 \pm 0.35% and 0.13 \pm 0.13%, respectively.

3.2. Enzymatic activity

Catalase activity was detected in all isolates. Similar results were also observed by Bautista-Gallego et al. (2011) and Hernández et al. (2007), in most yeasts isolated during fermentation of table olives. The ability of the yeast to produce this enzyme can be of great interest, since it is linked to the antioxidant capacity (Chen et al., 2010). In relation to lipase, *C. tropicalis* 3A, *C. boidinii* 32A and *G. reessii* 33A did not show any activity, while *P. manshurica* 2A and *S. cerevisiae* 15A and 15B showed high activity. The protease and xylanase activities were quite weak or null in most of the isolates assayed; however, lipase and protease can promote the probiotic potential (Syal & Vohra, 2013), since they contribute to the improvement of digestion of lipids and proteins in the gastrointestinal tract. Amylase is responsible for the hydrolysis of polysaccharides by facilitating the breakdown of starch and glycogen in the human diet, thus contributing to the digestive system (Motta, 2000). Nevertheless, only two strains (*C. boidinii* 32A and *G. reessii* 33A) showed activity of this enzyme. Regarding β -glucosidase, strong activity was detected in *R. glutinis* 27A and *R. graminis* 20A, while moderate activity was observed for *C. norvegica* 7A, *P. guilliermondii* 25A and *G. reessii* 33A. This enzyme affects the sensory characteristics of table olives, since β -glucosidase is involved in the biodegradation of polyphenols in fruit, reducing the oleuropein content, which is responsible for the bitterness of fruit (Arroyo-López et al., 2012).

3.3. Antioxidant activity

Table 2 shows the DPPH free radical scavenging activity for each yeast strain. The highest antioxidant activities were obtained for *S. cerevisiae* 15A (16.47 \pm 3.44%) and 15B (18.34 \pm 2.24%), similar to the reference strain *S. boulardii* (16.22 \pm 3.50%), while the lowest values were obtained for *C. boidinii* 32A (2.55 \pm 3.45%) and 37A (3.52 \pm 0.88%). Gil-Rodríguez et al. (2015), in a study with yeasts isolated from different food and beverages, obtained higher values for different strains of *S. cerevisiae*, ranging from 19.74 \pm 0.12% (IFI-88) to 56.97 \pm 0.30% (IFI-279), using a similar method to the present work. It is believed that the antioxidant activity of yeast is mainly due to the high content of (1,3)- β -D-glucan and other β -glucans found in the cell

wall (Abbas, 2006; Jaehrig, Rohn, Kroh, Fleischer, & Kurz, 2007), to other cellular compounds and due to the presence of antioxidant enzymes such as superoxide dismutase, glutathione peroxidase and catalase (Chen et al., 2010).

3.4. Autoaggregation capacity

Probiotic microorganisms, beyond resisting the conditions of the human gastrointestinal tract, should have good adherence to epithelial cells to be able to colonize and thus exert its beneficial properties. The autoaggregation ability is reflected in the ability of microorganisms to join together, forming cellular aggregates, thereby favoring gastrointestinal colonization (García-Cayuela et al., 2014). This ability is an important prerequisite for the colonization of probiotic strain in gastrointestinal tract, preventing their immediate elimination by the peristaltic movements and allowing a competitive advantage among existing microorganisms (Syal & Vohra, 2013).

The autoaggregation abilities of the assayed yeasts, determined at different time intervals, are presented in Table 2. At 2 h of incubation at 37 °C, the strains exhibited high variability in autoaggregation percentages, ranging between 2.86 \pm 1.18% (*R. glutinis* 27A) and 74.53 \pm 4.23% (*C. tropicalis* 1A). *C. tropicalis* showed autoaggregation percentage significantly higher ($p < 0.05$) than the other strains. Higher autoaggregation capacity of this strain was also reported by Syal and Vohra (2013), after 3 h of incubation (88.48 \pm 0.33%). After 4 h of incubation, the strains *S. cerevisiae* 15A and 15B, *C. norvegica* 7A, *C. boidinii* 37A and *P. manshurica* 2A, significantly increased their ability to form cellular aggregates, increasing continuously up to 24 h. The highest percentages of autoaggregation were determined after 24 h of incubation and ranged from 20.11 \pm 4.20% (*R. graminis* 20A) to 93.94 \pm 4.73% (*C. tropicalis* 1A). The variability in these results indicates that the autoaggregation ability is strongly strain dependent, as suggested by Gil-Rodríguez et al. (2015). Our autoaggregation results obtained at 24 h for *S. cerevisiae* 15A and 15B (91.71 \pm 5.58% and 74.33 \pm 6.93%, respectively) were similar, under the same condition to those values reported by Gil-Rodríguez et al. (2015) for *S. cerevisiae* IFI-87 (83.3 \pm 0.1%) and IFI-244 (99.8 \pm 5.74%), isolated from food and beverages. Syal and Vohra (2013) reported values of 100% after 20 h of incubation for *P. manshurica* and *S. cerevisiae* species.

3.5. Antimicrobial activity

Among the yeast strains studied, only two strains (*C. norvegica* 7A and *G. reessii* 34A) showed antimicrobial activity against *C. neoformans* (data not shown). Indeed, the strains *G. reessii* 34A and *C. norvegica* 7A showed inhibition halos of 2.7 and 1.5 cm, respectively. It should be noted that the antifungal agent used as a positive control (amphotericin B) showed a slightly lower inhibition halo (2.2 cm diameter) compared to the strain *G. reessii* 34A, suggesting that this strain has a strong antimicrobial activity against this microorganism. Antimicrobial activity

Table 2Antioxidant activity (%) and autoaggregation capacity (%) of the assayed yeasts strains after 2 h, 4 h and 24 h (mean \pm standard deviation, n = 3).

Isolates	Species	DPPH scavenging activity (%)	Autoaggregation (%)		
			2 h	4 h	24 h
1A	<i>C. tropicalis</i>	8.48 \pm 2.16 ^b	74.53 \pm 4.23 ^{A,a}	77.54 \pm 4.50 ^{A,a}	93.94 \pm 4.73 ^{B,a}
3A	<i>C. tropicalis</i>	7.85 \pm 0.76 ^b	65.5 \pm 0.24 ^{A,a,b}	71.29 \pm 1.84 ^{A,a,b}	78.37 \pm 3.57 ^{B,b,c}
29A	<i>P. membranifaciens</i>	ND	48.16 \pm 7.71 ^{A,c,d}	47.11 \pm 4.22 ^{A,c,d}	76.13 \pm 5.46 ^{B,c}
3B	<i>P. membranifaciens</i>	ND	54.71 \pm 8.56 ^{A,b,c}	65.60 \pm 4.07 ^{A,B,a,b}	77.08 \pm 1.70 ^{B,c}
15B	<i>S. cerevisiae</i>	18.34 \pm 2.24 ^a	53.73 \pm 6.67 ^{A,b,c}	71.23 \pm 3.78 ^{B,a,b}	74.33 \pm 6.93 ^{B,c}
15A	<i>S. cerevisiae</i>	16.47 \pm 3.44 ^a	59.97 \pm 3.09 ^{A,b}	71.92 \pm 6.30 ^{B,a}	91.71 \pm 5.58 ^{C,a,b,c}
7A	<i>C. norvegica</i>	6.57 \pm 0.85 ^b	38.27 \pm 0.30 ^{A,d,e}	59.21 \pm 5.24 ^{B,b,c}	82.72 \pm 7.14 ^{C,a,b,c}
8A	<i>D. hansenii</i>	4.41 \pm 1.95 ^b	31.49 \pm 3.55 ^{A,e,f}	30.83 \pm 3.27 ^{A,e,f}	34.37 \pm 10.80 ^{A,d}
20A	<i>R. graminis</i>	ND	3.35 \pm 0.71 ^{A,h}	4.05 \pm 1.42 ^{A,h}	20.11 \pm 4.20 ^{B,d}
37A	<i>C. boidinii</i>	3.52 \pm 0.88 ^b	26.46 \pm 2.95 ^{A,e,f}	41.82 \pm 4.16 ^{B,d,e}	76.58 \pm 7.90 ^{C,c}
32A	<i>C. boidinii</i>	2.55 \pm 3.45 ^b	8.96 \pm 4.53 ^{A,h}	13.58 \pm 6.40 ^{B,g,h}	25.36 \pm 3.69 ^{B,d}
25A	<i>P. guilliermondii</i>	7.92 \pm 2.15 ^b	25.05 \pm 5.13 ^{A,e,f,g}	30.66 \pm 6.66 ^{A,e,f}	73.66 \pm 2.65 ^{B,c}
27A	<i>R. glutinis</i>	ND	2.86 \pm 1.18 ^{A,h}	5.12 \pm 1.88 ^{A,h}	23.67 \pm 8.68 ^{B,d}
2A	<i>P. manshurica</i>	ND	12.82 \pm 2.00 ^{A,g,h}	24.75 \pm 5.05 ^{B,f,g}	76.86 \pm 1.46 ^{C,c}
33A	<i>G. reessii</i>	8.14 \pm 1.72 ^b	22.01 \pm 4.95 ^{A,f,g}	29.21 \pm 5.51 ^{A,e,f}	70.50 \pm 3.10 ^{B,c}
34A	<i>G. reessii</i>	ND	34.86 \pm 4.27 ^{A,e,f}	30.55 \pm 4.88 ^{A,e,f}	34.48 \pm 4.16 ^{A,d}
Sb	<i>S. boulardii</i>	16.22 \pm 3.50 ^a	25.50 \pm 5.81 ^{A,f}	34.75 \pm 2.95 ^{A,d,e,f}	77.05 \pm 5.74 ^{B,c}

Different lowercase letters in the same column indicate statistically differences ($p < 0.05$). For the autoaggregation results, different uppercase letters in the same line indicate significant differences ($p < 0.05$) by Tukey's test. ND – not determined.

observed in these strains could be related to the ability of yeasts that can produce and deliver a substance of proteinaceous nature known as mycocin or killer toxin, which is lethal for sensitive yeasts of the same species or of other genera (Banjara, Nickerson, Suhr, & Hallen-Adams, 2016; Gil-Rodríguez et al., 2015).

Concerning the probiotic reference strain (*S. boulardii*), it has not been demonstrated antimicrobial activity against the tested pathogens. Nevertheless, Czerucka, Piche, and Rampal (2007) observed antimicrobial effect of this yeast for various enteric pathogens such as *Clostridium difficile*, *Vibrio cholerae*, *Salmonella*, *Shigella* and *E. coli*.

3.6. Principal Component Analysis (PCA)

Principal Component Analysis (PCA) was used in order to analyze simultaneously all the relevant variables. The variables used corresponded to the results of protease (Prot), β -glucosidase (B-gluc), xylanase (Xil), lipase (Lip), antimicrobial activity (AM), antioxidant activity (AA), autoaggregation (Ag2h, Ag4h and Ag24h), survival of gastric (SG) and pancreatic (SP) digestions, and total survival (ST).

The biplot of the principal components (Fig. 1) showed the existence of two major factors that explained 54.21% of the total variance of the original variables (34.29% for Factor 1 and 19.92% for Factor 2). The first component was mainly related to lipase, antioxidant activity, autoaggregation at 2, 4 and 24 h, gastric survival, and total survival. The second component (Factor 2) was mainly related to xylanase, β -glucosidase, antimicrobial activity and pancreatic survival (Fig. 1). The assayed yeast strains could be classified into four different groups. The first group included the strain *C. norvegica* 7A which showed the highest antimicrobial activity. The second group included the strain *P. guilliermondii* 25A, which presented the highest total survival. The third group included the strains *D. hansenii* 8A, *C. boidinii* 37A and *C. tropicalis* 3A. *D. hansenii* 8A was the one with the lowest values for the antioxidant activity and autoaggregation ability, while *C. tropicalis* 3A showed the highest survival to gastric digestion. The remaining strains were included in the fourth group.

4. Conclusions

In this study we isolated yeast strains that could be used as new probiotic strains. *S. cerevisiae* showed lipase activity, high ability to grow at the host temperature and high antioxidant capacity, *S. cerevisiae* 15A showed higher autoaggregation percentage than the reference

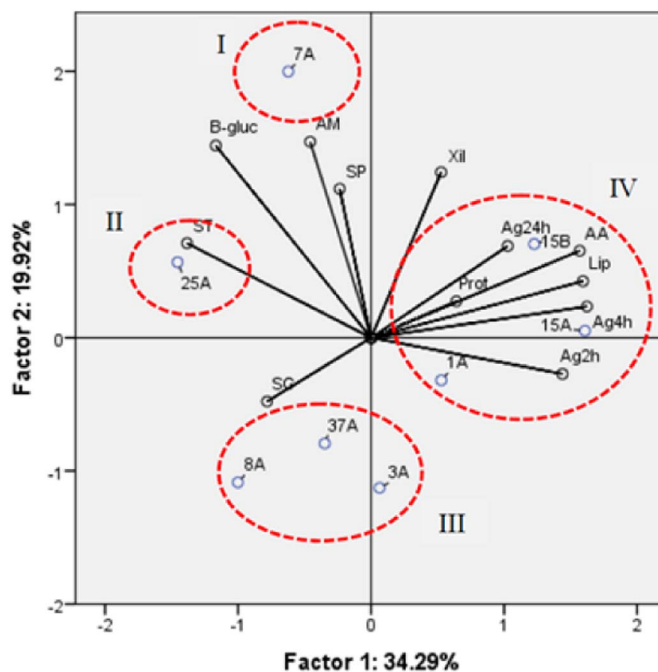


Fig. 1. Loadings and scores plot resulting from the Principal Component Analysis applied to the yeast strains, considering the following variables: AA: antioxidant activity; Ag2h, Ag4h and Ag24h: autoaggregation at 2, 4 and 24 h, respectively; Survival of gastric (SG), pancreatic (SP), and total digestions (ST); Prot: protease; Xil: xylanase; B-gluc: β -glucosidase; AM: amylase; and Lip: lipase.

strain. *P. guilliermondii* and *C. norvegica* exhibited ability to survive to the human gastrointestinal tract digestion. Our results also demonstrated that *C. norvegica* have antimicrobial activity against *C. neoformans*. Nevertheless, it would be interesting to perform additional tests to these strains to better understand the mechanisms involved, taking into account their role in table olives fermentation and table olives characteristics, as well as to evaluate other probiotic properties, including hydrophobicity, antibiotic resistance, and adhesiveness to Caco-2 enterocytes. Furthermore, *in vivo* testing using animal models should be developed in order to ascertain the benefits observed *in vitro*.

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