

# Interleukin 6 Causes Growth Impairment in Transgenic Mice through a Decrease in Insulin-like Growth Factor-I

## A Model for Stunted Growth in Children with Chronic Inflammation

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### Abstract

Stunted growth is a major complication of chronic inflammation and recurrent infections in children. Systemic juvenile rheumatoid arthritis is a chronic inflammatory disorder characterized by markedly elevated circulating levels of IL-6 and stunted growth. In this study we found that NSE/hIL-6 transgenic mouse lines expressing high levels of circulating IL-6 since early after birth presented a reduced growth rate that led to mice 50–70% the size of nontransgenic littermates. Administration of a monoclonal antibody to the murine IL-6 receptor partially reverted the growth defect. In NSE/hIL-6 transgenic mice, circulating IGF-I levels were significantly lower than those of nontransgenic littermates; on the contrary, the distribution of growth hormone pituitary cells, as well as circulating growth hormone levels, were normal. Treatment of nontransgenic mice of the same strain with IL-6 resulted in a significant decrease in IGF-I levels. Moreover, in patients with systemic juvenile rheumatoid arthritis, circulating IL-6 levels were negatively correlated with IGF-I levels. Our findings suggest that IL-6-mediated decrease in IGF-I production represents a major mechanism by which chronic inflammation affects growth. (*J. Clin. Invest.* 1997; 99:643–650.) Key words: interleukin 6 • insulin-like growth factor-I • growth disorders • juvenile rheumatoid arthritis

### Introduction

Stunted growth is one of the major complications of chronic inflammation and infection in children. Although several causes, including poor nutrition, immobilization, and malabsorption, have been hypothesized, the mechanism responsible for growth impairment in these conditions is unknown (1).

Systemic juvenile rheumatoid arthritis (s-JRA)<sup>1</sup> is a chronic

inflammatory disease associated with stunted growth (2, 3). Growth impairment in s-JRA occurs during periods of disease activity with subsequent normalization of growth rate during remission (4, 5). Growth hormone (GH) production is normal in patients with s-JRA (6, 7), while levels of insulin-like growth factor-I (IGF-I) are decreased (7–9). IGF-I is produced by the liver in response to GH, mediates GH effects in several peripheral organs including muscles and bones (10), and has a pivotal role in postnatal growth as demonstrated by the observation that mice carrying a null mutation of the IGF-I gene showed markedly decreased postnatal growth (11). In s-JRA, laboratory evidence of inflammation is prominent (3), and interleukin 6 (IL-6) appears to be very relevant to the inflammatory process of s-JRA; circulating and synovial levels of IL-6 are markedly elevated in patients with active s-JRA, are significantly higher than in other adult or juvenile chronic arthritides, and are related to clinical and laboratory parameters of disease activity; IL-6 appears to explain several clinical and laboratory features of the disease (12–17).

In this study we have found that the NSE/hIL-6 transgenic murine lines with high levels of circulating IL-6 since early phases of life show a growth defective phenotype; we demonstrate that IL-6 is associated with a growth defect and causes a decrease in IGF-I levels, without affecting GH production. Our findings show that sustained production of IL-6 may represent the mechanism by which chronic inflammation affects growth.

### Methods

**Animals and treatments.** The NSE/hIL-6 construct carries the rat neurospecific enolase (NSE) promoter driving the expression of human IL-6 cDNA. The generation of NSE/hIL-6 mice has been described recently (18). Overexpression of hIL-6 in neuronal tissue results in reactive astrocytosis and in an increase in ramified microglial cells, but these mice do not show histological or behavioral signs of neuron damage (18). Transgenic animals were identified by PCR analysis of DNA extracted from a tail segment, as described (18). Mice were maintained in standard conditions under a 12-h light–dark cycle and were provided irradiated food (4RF21, Mucedola; Settimo Milanese, Milan, Italy) and chlorinated water ad libitum. Procedures involving animals and their care were conducted in conformity with national and international laws and policies (EEC Council Directive 86/609, OJ L 358, 1, December 12, 1987; Italian Legislative Decree 116/92, Gazzetta Ufficiale della Repubblica Italiana n. 40, February 18, 1992; National Institutes of Health Guide for the Care and Use of Laboratory Animals, publication No. 85–23, 1985). To evaluate the effect of the inhibition of hIL-6 action, NSE/hIL-6 mice were injected subcutaneously at days 2, 4, 7, 11, 14, and 20 after birth (day 0) with the neutralizing rat monoclonal antibody 15A7 to the murine IL-6 receptor (19) at doses of 90  $\mu$ g/mouse at day 2, 150  $\mu$ g/mouse at day 4, and of 300  $\mu$ g/mouse at day 7 and thereafter. The antibody was pre-

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1. Abbreviations used in this paper: GH, growth hormone; NSE, neurospecific enolase; s-JRA, systemic juvenile rheumatoid arthritis; TSH, thyroid stimulating hormone.

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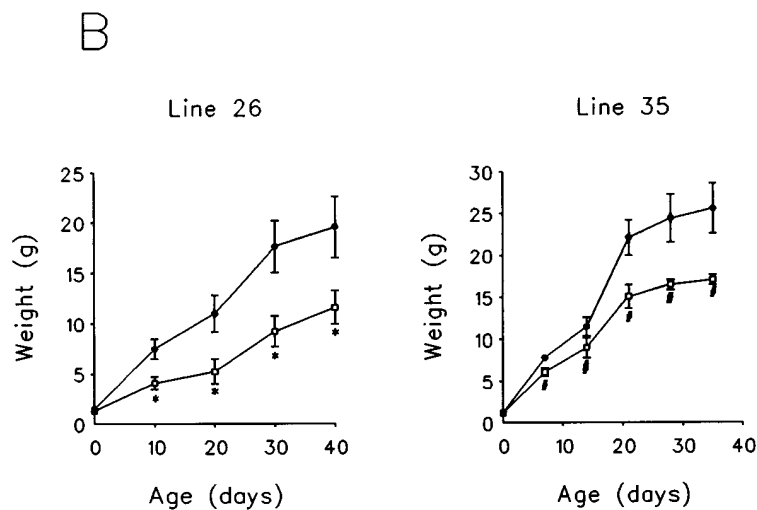
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cipitated from the hybridoma cell medium in saturating ammonium sulphate, dialyzed against 50 mM Tris/HCl, pH 7.5, and loaded on a protein G–Sepharose resin (cat. No. 17-0618-02; Pharmacia, Brussels, Belgium), equilibrated with the same buffer. Elution was performed with 100 mM sodium citrate/citric acid buffer, pH 3.0, and the fractions collected were immediately neutralized with 3 M Tris/HCl, pH 8.9. Before injection the antibody was dialyzed against sterile pyrogen-free saline solution. Control mice were treated with sterile pyrogen-free saline solution. To evaluate the effect of administration of IL-6 to CB6F1 (C57Bl6xBalb/C) mice, recombinant human IL-6 (rhIL-6), obtained as described (20), resuspended in sterile pyrogen-free saline solution, was administered intraperitoneally at a dose of 10  $\mu$ g/mouse to 3-wk-old animals; control mice were injected intraperitoneally with sterile pyrogen-free saline solution. To measure food intake and body weight, food and mice were weighed every day at the same hour. Food intake per gram of body weight was calculated by dividing the amount of food consumed over a 24-h period for the mean mouse weight at the end of the same period. Hematic glucose was determined using the Accutrend/GC apparatus and Strips Accutrend Glucose (Boehringer-Mannheim, Mannheim, Germany). The pituitaries were carefully dissected from the cranial base and fixed in 2% paraformaldehyde for 1 h. The embedding and immunostaining were performed as described (21). Antisera specific for mouse GH and for rat thyroid stimulating hormone (TSH) were kindly provided by the National Hormone and Pituitary Program at the National Institute of Health (Bethesda, MD); the secondary antibodies labeled

with peroxidase were purchased from Jackson ImmunoResearch Laboratories, Inc. (West Grove, PA).

**Patients.** 21 patients (mean age 6.5 yr, age range 2–17 yr) fulfilling the American College of Rheumatology (formerly American Rheumatism Association) criteria for the diagnosis of s-JRA (22) were included in the study. All patients presented active disease at time of sampling as defined by the presence of synovitis on examination and were receiving nonsteroidal antiinflammatory drugs; in addition approximately half of them were also treated with low-dose steroids (on alternate day regimen in the majority); two-thirds of the patients were also receiving methotrexate. Permission for drawing of extra blood during routine venipuncture was obtained from parents of all children. Since marked changes in circulating IL-6 levels occurs during the febrile peak (12, 16), peripheral blood samples were collected during the morning hours, when all patients were afebrile.

**Measurement of IGF-I, IL-6, IL-1 $\beta$ , and TNF- $\alpha$ .** IGF-I was measured using a commercially available radioimmunoassay, which recognizes both murine and human IGF-I, according to the instructions provided by the manufacturer (Nichols, San Juan Capistrano, CA), after acid-ethanol extraction of plasma samples (anticoagulated with EDTA, final concentrations 5 mg/ml). IGF-I levels in patients with s-JRA were compared with normal values for age as determined by the manufacturer in 944 healthy controls. In patients with s-JRA, serum IL-6 levels were measured with a hybridoma growth factor assay using B9 cells as described previously (12); the hybridoma growth factor activity in sera of patients with s-JRA was abolished by the addi-



**Figure 1.** (A) Reduced size of a 12-d-old NSE/hIL-6 mouse of line 26 (arrow) compared with an age- and sex-matched nontransgenic littermate. (B) Growth curves of transgenic (open squares) and nontransgenic (closed circles) NSE/hIL-6 mice of lines 26 and 35. Numbers of each group of mice are as described in the legend to Table I. Results are shown as means  $\pm$  standard deviations (represented by the vertical bars). \* $P < 0.001$  and  $^{\#}P < 0.005$  versus nontransgenic littermates.

Table I. Growth Rate of Transgenic and Nontransgenic NSE/hIL-6 Mice of Lines 26 and 35 in the Indicated Age Periods

Growth rate (mean gram increase in body weight per day)					
Line 26					
Age period (d)	0–10	11–20	21–30	31–40	
Transgenic	0.28±0.07	0.13±0.10	0.37±0.10	0.20±0.12	
Wild-type	0.59±0.10	0.36±0.10	0.62±0.17	0.22±0.13	
<i>P</i>	< 0.0001	0.0001	0.0002	0.34	
Line 35					
Age period (d)	0–7	8–14	15–21	22–28	29–35
Transgenic	0.71±0.06	0.41±0.07	0.87±0.14	0.22±0.08	0.09±0.05
Wild-type	0.94±0.08	0.53±0.10	1.52±0.38	0.33±0.09	0.18±0.05
<i>P</i>	0.0034	0.027	0.008	0.063	0.010

The weight was determined at the same hour of the day once every 10 d for mice of line 26 and once a week for mice of line 35 (line 26: transgenics *n* = 14, wild-type *n* = 17; line 35: transgenics *n* = 7, wild-type *n* = 6). Results are shown as means±standard deviations.

tion of a goat antiserum to hIL-6 (not shown). In mice, IL-6 levels were determined using an immunoassay specific for biologically active human IL-6 (Boehringer-Mannheim). In patients with s-JRA, TNF- $\alpha$  and IL-1 $\beta$  were measured in plasma samples (anticoagulated with EDTA, as above) using two commercially available immunoassays (TNF- $\alpha$  from Boehringer-Mannheim; IL-1 $\beta$  from Immunotech, Marseille, France).

**Statistical analysis.** Results were analyzed using the Mann-Whitney U test for unpaired samples and the Spearman correlation coefficient, as appropriate. A *P* value < 0.05 was considered significant. A Z score for plasma IGF-I levels of patients with s-JRA was calculated according to the following formula: log (patient value) – log (mean value for the age control group)/log (standard deviations of the age control group).

## Results

**Growth defect in NSE/hIL-6 mice.** The generation of NSE/hIL-6 mice has been reported recently (18). Four of the seven transgenic founders generated (mice 15, 22, 26, and 35) transmitted the transgene to the progeny and four stable lines were established. No evident differences in body weight were revealed between transgenic and wild-type mice of the four lines at birth. After birth, while no differences were noted in the growth rate of transgenics of lines 15 and 22 compared with nontransgenic littermates (not shown), transgenic mice of lines 26 and 35 presented a reduced growth rate that led to mice 50–70% the size of the age-matched littermates (Fig. 1 A). Growth rates and growth curves of lines 26 and 35 are shown in Table I and Fig. 1 B, respectively.

Subsequent analysis showed that lines 26 and 35 had high levels of circulating hIL-6, detectable already 4 d after birth, while transgenics of lines 15 and 22 had undetectable levels (Table II). Moreover, Northern blot analysis showed that the levels of hIL-6 expression in the brain were similar in the four lines; however, while in lines 15 and 22 the expression of the transgene was confined to the central nervous system, in lines 26 and 35 hIL-6 expression was detectable in heart and lung; no hIL-6 expression was detected in liver, spleen, or kidney (18).

To verify whether the growth defect was due to the elevated levels of circulating IL-6, we tried to inhibit the action of the cytokine in transgenic mice of line 26 by injecting the monoclonal antibody 15A7 neutralizing the murine IL-6 receptor,

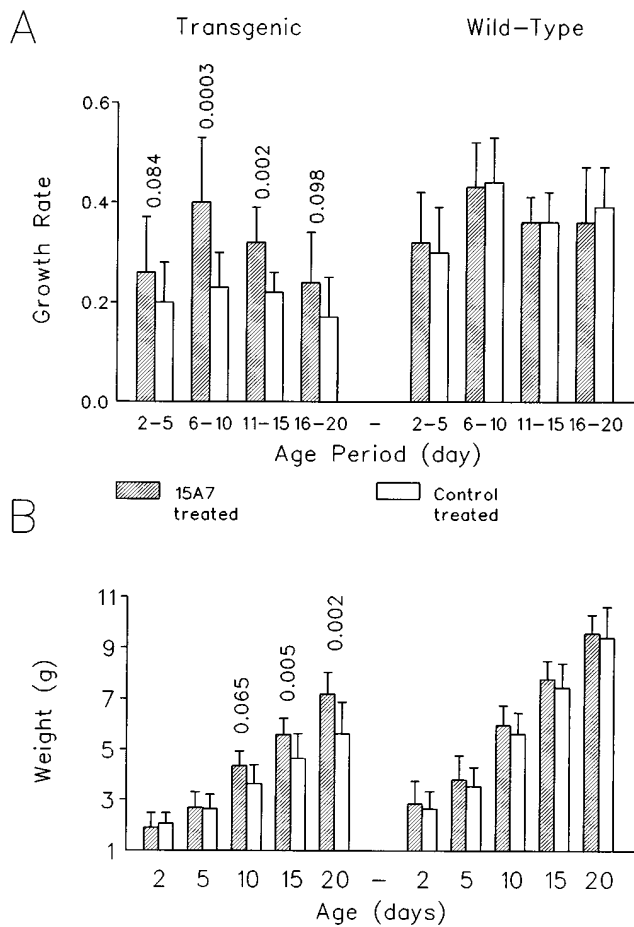
starting at day 2 after birth. Neutralization of IL-6 activities resulted in a significant improvement in the growth rate, that was particularly evident from days 6 to 15 (Fig. 2 A). The nonstatistically significant effect on growth rate observed from days 15 to 20 may be due to some endogenous production of antibodies directed against the rat monoclonal antibody, and/or to the insufficiency of the dose administered relative to the increase in body weight (i.e., a constant dose of 300  $\mu$ g/mice was administered at days 7, 11, 14). As shown in Fig. 2 B, the improvement in growth rate resulted in a partial, but significant, correction of the growth defect, as indicated by the statistically significant differences in weight at days 15 (*P* = 0.005) and 20 (*P* = 0.002) of age between transgenic mice treated with the monoclonal antibody 15A7 (weight at day 20: 7.19±0.84 g) and the control-treated transgenic animals (weight at day 20: 5.63±1.25 g). In nontransgenic mice no significant differences were found between 15A7-treated (weight at day 20: 9.56±0.74 g) and control-treated animals (weight at day 20: 9.40±1.21 g) (Fig. 2, A and B). These data indicate that in NSE/hIL-6 mice the growth defective phenotype is related to the elevated levels of IL-6.

**Food intake and hematic glucose in NSE/hIL-6 mice.** Reduction of food intake and hypoglycemia are common host re-

Table II. Serum hIL-6 Levels in NSE/hIL-6 Transgenic Mice at Different Ages

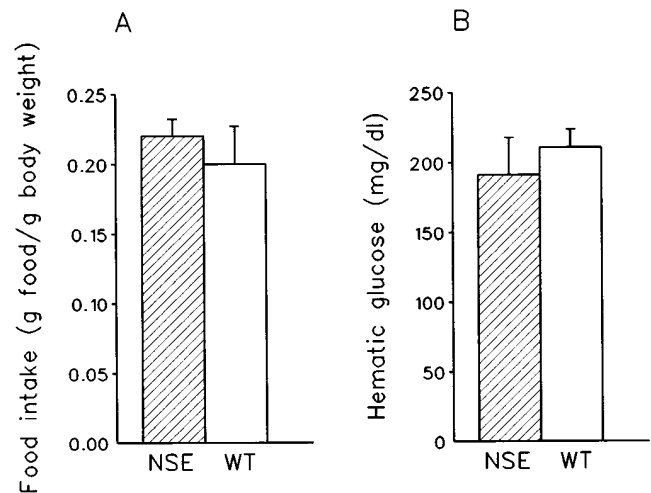
	hIL-6 (ng/ml)			
	Line 15	Line 22	Line 26	Line 35
Day 4	ND	ND	5.10±0.84 ( <i>n</i> = 3)	3.90±1.90 ( <i>n</i> = 2)
Day 11	ND	ND	3.52±1.75 ( <i>n</i> = 4)	ND
Day 20	< 0.06	< 0.06	15.0±5.0 ( <i>n</i> = 5)	6.0±0.80 ( <i>n</i> = 6)

Mice of lines 15 and 22 have normal growth, mice of lines 26 and 35 show a growth defect. In nontransgenic mice serum hIL-6 levels were always below the detection limit (i.e., 0.06 ng/ml). Results are shown as means±standard deviations. ND, not determined.



**Figure 2.** Effect of the neutralization of IL-6 activity on the growth of NSE/hIL-6 transgenic or wild-type mice of line 26. *A* shows the growth rate (mean gram increase in body weight per day) in the indicated age periods. *B* shows the weight at the indicated ages. Transgenic mice are shown on the left, and wild-type mice on the right. Significance levels (*P* values) of the differences of 15A7-treated animals versus corresponding control-treated animals are shown when *P* < 0.1. Mice were treated with the monoclonal antibody 15A7 neutralizing the murine IL-6 receptor (shaded columns) or with saline (open columns), as described in Methods (control-treated transgenic *n* = 14; 15A7-treated transgenic *n* = 14; control-treated wild-type *n* = 20; 15A7-treated wild-type *n* = 19). The results are shown as means  $\pm$  standard deviations (represented by the vertical bars).

sponses to infection and inflammation and there is growing evidence that these phenomena result in part from the release of proinflammatory cytokines (23–26). Moreover, high levels of circulating IL-6 have been revealed during starvation in patients with anorexia nervosa (27). Therefore, to determine whether the reduced growth rate of NSE/hIL-6 mice could be due to a behavioral disorder leading to a reduction of food intake with subsequent hypoglycemia, we measured the food intake of age-matched transgenic and wild-type mice of line 26 over a period of 7 d starting at 4 wk of age. Although the amount of food consumed daily was, as expected, lower in the small size NSE/hIL-6 transgenics than in wild-type littermates (not shown), the food intake per gram of body weight was comparable in the two groups (Fig. 3 *A*). Moreover, no differences were found in the levels of hematic glucose (Fig. 3 *B*).



**Figure 3.** (*A*) Food intake in NSE/hIL-6 transgenic mice of line 26 (*n* = 5) (shaded bars) and wild-type littermates (*n* = 5) (open bars) expressed as daily food intake per gram of body weight. The two groups of mice were caged separately and the amount of food was weighed once a day at 9 a.m.; the results are the means of 7 d of measurements  $\pm$  standard deviations. (*B*) Hematic glucose levels in 8-wk-old NSE/hIL-6 transgenic mice of line 26 (*n* = 5) (shaded bars) and in wild-type littermates (*n* = 5) (open bars); results are shown as means  $\pm$  standard deviations.

These results indicated that the growth defect in NSE/hIL-6 mice is not caused by a nutritional disorder.

**Decreased *in vivo* levels of IGF-I in NSE/hIL-6 mice.** The pituitary hormones GH and TSH play a major role in somatic growth; to investigate the possible effects of IL-6 overexpression on pituitary functions, we evaluated the distribution of GH- and TSH-producing pituitary cells in mice of lines 26 and 35. Immunocytochemical analysis did not show differences between transgenic and wild-type littermates (Fig. 4). In addition, circulating levels of GH, as well as of T4, were comparable between transgenic and wild-type littermates (not shown).

Since IGF-I mediates the great majority of the peripheral effects of GH (10) and plays a pivotal role in postnatal growth (11), and since in s-JRA the growth defect is associated with low levels of IGF-I (7–9), we measured circulating IGF-I levels in NSE/hIL-6 mice of lines 26, 35, and 22. As shown in Table III, in transgenic mice of lines 26 and 35, IGF-I levels were significantly lower than in the corresponding age-matched wild-type littermates, while no differences between transgenics and nontransgenics were found for line 22 with undetectable circulating IL-6 and normal growth. Further supporting the role of the decrease in IGF-I in the growth defect of line 26, a positive correlation between IGF-I levels and body weight at day 20 after birth was found both in transgenic ( $R_s = 0.743$ ,  $P = 0.005$ ) and nontransgenic ( $R_s = 0.788$ ,  $P < 0.001$ ) mice of line 26. These results show that in both lines 26 and 35 with stunted growth, elevated circulating levels of IL-6 are associated with a decrease in IGF-I levels to values that are < 50% of those present in the corresponding nontransgenic animals.

**IL-6-induced decrease in *in vivo* IGF-I levels in CB6F1 mice.** To verify the relationship between elevated levels of IL-6 and the decrease in circulating IGF-I, we injected recombinant hIL-6 in mice of the same strain of the NSE/hIL-6 transgenics. As shown in Fig. 5 *A*, treatment of CB6F1 mice with two intra-

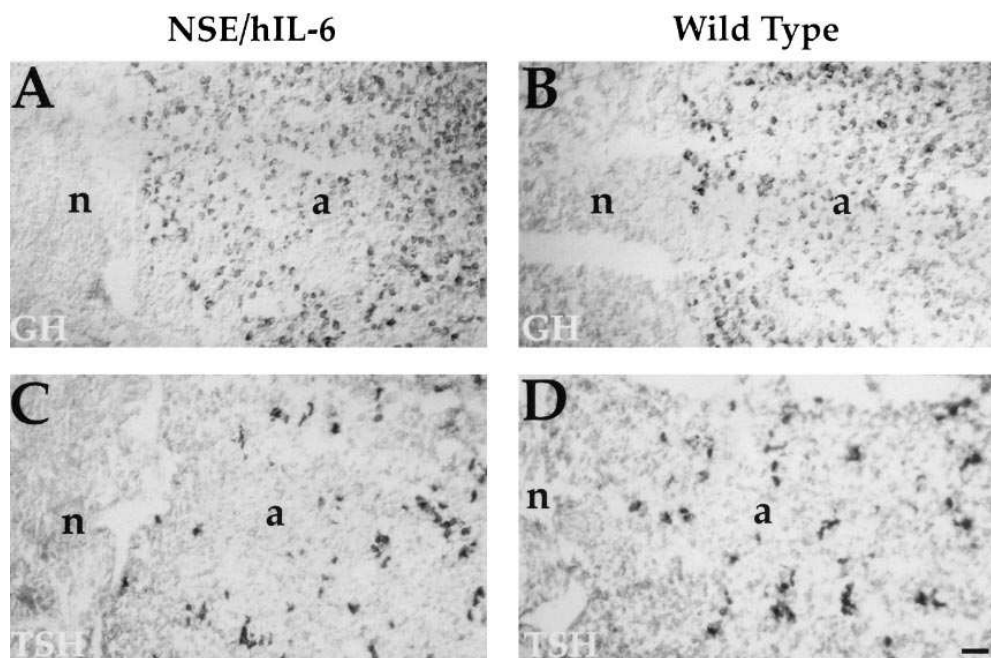


Figure 4. Distribution of GH- or TSH-producing pituitary cells in NSE/hIL-6 mice. Frontal sections of pituitaries from NSE/hIL-6 (A and C) and wild-type littermates (B and D) were immunostained with antibodies specific for GH (A and B) or TSH (C and D). *n*, neurohypophysis; *a*, adenohypophysis. Bar = 25  $\mu$ m.

peritoneal administrations of recombinant hIL-6 at 12-h intervals resulted in a significant decrease in IGF-I levels 24 h after the first treatment compared with untreated or saline-treated mice. Although a modest, albeit not significant, reduction in weight gain in the 2 d subsequent to hIL-6 administration was observed (weight gain hIL-6 treated:  $0.24 \pm 0.39$  g; weight gain control-treated:  $0.53 \pm 0.38$  g;  $P = 0.16$ ), this did not result in a significant effect on the body weight of CB6F1 mice for the entire length of the period examined (Fig. 5 B). The absence of a

significant effect of this treatment on body weight suggests that prolonged overproduction of IL-6 is necessary to induce defective growth. IGF-I levels are strongly reduced upon fasting both in human and rats (28). To verify if our protocol of IL-6 administration affected food intake and therefore influenced only indirectly IGF-I levels, we measured food intake in mice starting 2 d before treatment and lasting 6 d after. A reduction in food intake was observed in the IL-6-treated mice, but only starting 2 d after treatment, i.e., 24 h after the determination of IGF-I levels (Fig. 5 C). Based on these results and on the normal food intake and glycemia of NSE/hIL-6 mice, we concluded that IL-6 affects the circulating levels of IGF-I by mechanisms that are distinct from anorexia.

Table III. Circulating IGF-I Levels in Transgenic and Nontransgenic NSE/hIL-6 Mice of Lines 26 and 35 (Elevated Circulating hIL-6 and Growth Defect) and of Line 22 (No Circulating hIL-6 and Normal Growth), at the Indicated Ages

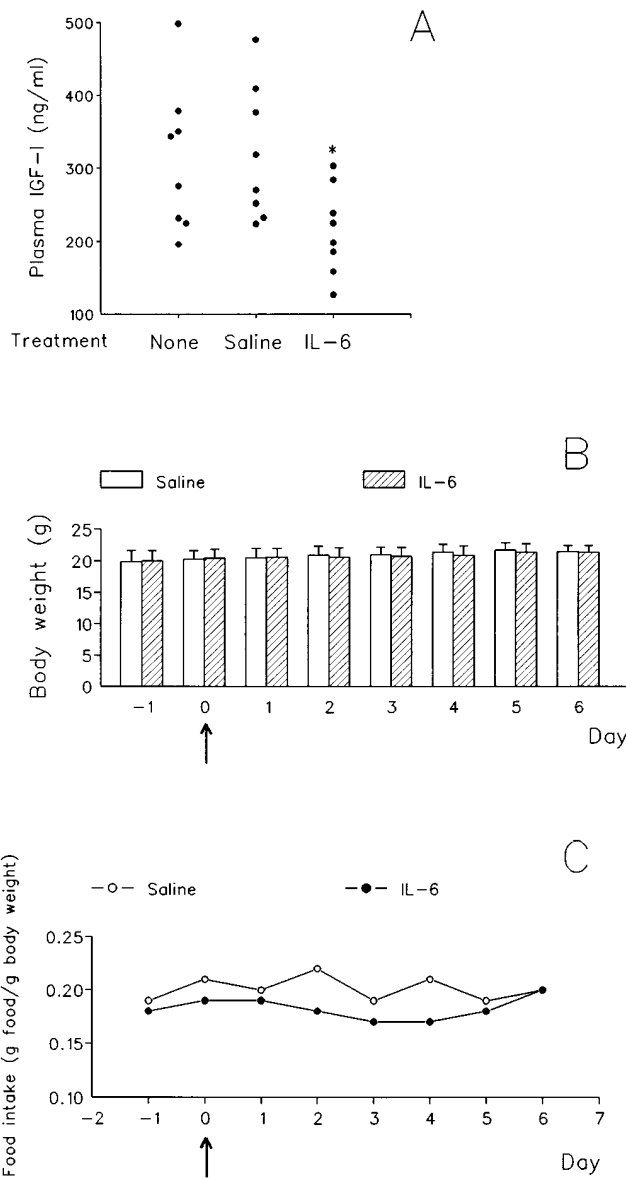
	IGF-I (ng/ml)		
	Age	11 d	20 d
Line 26			
Transgenic		$108.7 \pm 42.5^*$ ( <i>n</i> = 3)	$124.6 \pm 40.6^\ddagger$ ( <i>n</i> = 11)
Wild-type		$215 \pm 328.3$ ( <i>n</i> = 3)	$271.9 \pm 63.0$ ( <i>n</i> = 14)
Line 35			
Transgenic		ND	$38.4 \pm 8.6^\ddagger$ ( <i>n</i> = 3)
Wild-type		ND	$145.6 \pm 58.7$ ( <i>n</i> = 7)
Line 22			
Transgenic		ND	$224.0 \pm 93.3$ ( <i>n</i> = 8)
Wild-type		ND	$223.0 \pm 66.2$ ( <i>n</i> = 7)

Results are shown as means  $\pm$  standard deviations. \* $P < 0.05$  and  $^\ddagger P < 0.005$  versus corresponding nontransgenic animals. ND, not determined.

*Negative correlation between IL-6 and IGF-I levels in patients with s-JRA.* In agreement with previous studies (7–9), we found that in patients with s-JRA plasma IGF-I levels were lower than the mean normal values for age, without any relation of IGF-I levels with steroid treatment (not shown). As shown in Fig. 6, IGF-I plasma levels were negatively correlated ( $R_s = -0.667$ ,  $P = 0.004$ ) with serum IL-6 levels. A significant negative correlation of IGF-I levels was also found with C-reactive protein concentrations ( $R_s = -0.499$ ,  $P = 0.02$ ). In contrast, IGF-I levels were not correlated with IL-1 $\beta$  or TNF- $\alpha$  levels ( $R_s = 0.125$ ,  $P > 0.1$ ;  $R_s = -0.243$ ,  $P > 0.1$ , respectively).

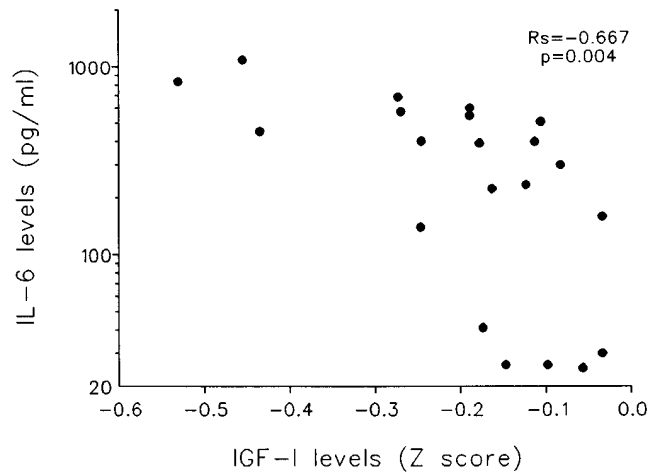
## Discussion

In this study we present the analysis of a transgenic mouse model that associates a growth defective phenotype with high circulating levels of IL-6 since birth. Several evidences demonstrate that the growth impairment present in NSE/hIL-6 mice does not result from an integration effect altering the expression of genes involved in growth: (a) the growth defect occurred in five of the seven founder mice generated (not shown) and in two different lines of transgenic offspring; (b) NSE/hIL-6 mice of lines 15 and 22, which have expression of



**Figure 5.** (A) Effect of the administration of recombinant hIL-6 to CB6F1 mice on circulating IGF-I levels. 3-wk-old CB6F1 mice (8 mice/group) were treated with two intraperitoneal injections of recombinant human IL-6 (10  $\mu$ g/mouse) or of sterile pyrogen-free saline or left untreated. Injections were administered at 12-h intervals (8 a.m. and 8 p.m., day 0). Blood was collected 12 h after the second injection (8 a.m., day 1) and circulating IGF-I was measured. \* $P = 0.03$  versus untreated or saline-treated mice. (B and C) Effect of IL-6 administration on body weight (B) and food intake, expressed as daily food intake per gram of body weight (C), in CB6F1 mice. 3-wk-old mice (8 mice/group, caged separately) were treated with saline or with recombinant human IL-6 as described in A; arrows indicate the time of treatment. Body and food weights were determined every day at 9 a.m. (at day 0 before the first administration of recombinant human IL-6).

the transgene confined to the CNS, grow with a normal rate, indicating that neither the presence of the transgene nor its expression in the CNS is per se sufficient to cause a growth defect; on the contrary, in NSE/hIL-6 mice of lines 26 and 35,



**Figure 6.** Negative correlation between serum IL-6 and plasma IGF-I levels in patients with s-JRA. To allow comparisons between values obtained from patients with different ages, plasma IGF-I levels are shown as Z scores, calculated as described in Methods. The Spearman correlation coefficient (Rs) and significance level (P) are shown.

which present growth impairment, hIL-6 mRNA expression is detected also in other organs (18) and this leaky expression, of unknown origin, is associated with measurable circulating IL-6 levels; and (c) the growth defect of NSE/hIL-6 mice can be partially corrected by neutralization of the murine IL-6 receptor, thus demonstrating that it is indeed due to IL-6 overexpression. In addition, other IL-6 transgenic mice show defective growth, including a neuron-, a keratinocyte-, and an airway epithelial cell-specific IL-6 transgenic (29–31), and although this aspect has never been described in detail, in some of them the defect has been reported since very early in life (29, 30). The fact that not all of the IL-6 transgenic mice described so far have growth impairment may be accounted for by different cell type and/or organ expression of IL-6, or by differential expression of the transgene with age. Indeed, in our experience, while in NSE/hIL-6 mice of lines 26 and 35 circulating hIL-6 in the range of nanograms per milliliter is detectable early after birth, MT-I/IL-6 mice, which carry the hIL-6 cDNA under the control of the mouse metallothionein-I promoter, do not show a growth defect and have circulating hIL-6 levels that increases markedly only after 2 mo of age (32).

The mechanism by which IL-6 causes growth defect appears to involve a decrease in circulating IGF-I levels. In NSE/hIL-6 mice of both lines 26 and 35, IGF-I levels are approximately half of those of the corresponding nontransgenic littermates; incidentally, in both transgenic and nontransgenic mice of line 26 we found a positive correlation between IGF-I levels and weight at day 20 after birth. Moreover, treatment of CB6F1 mice with recombinant IL-6 results in a decrease in IGF-I levels. Anorexia and hypoglycemia may be observed during acute and chronic inflammation and have been related to the action of inflammatory cytokines (33, 34); in turn, caloric and protein restrictions have been shown to cause a decrease in circulating IGF-I levels both in humans and rats (28). Our transgenic mice have a normal food intake, and a reduction of food intake in IL-6-treated CB6F1 mice occurs only 24 h later than the decrease in IGF-I, demonstrating that the de-

creased levels of IGF-I caused by elevated levels of IL-6 cannot be ascribed to nutritional disorders. NSE/hIL-6 mice of lines 26 and 35 have a normal distribution of GH-producing pituitary cells and normal GH levels, therefore demonstrating that the effect of IL-6 on IGF-I levels is not mediated indirectly via an effect on GH production. In addition, since the GH/IGF-I axis plays the major role in postnatal growth in the presence of normal thyroid function, it is noteworthy that NSE/hIL-6 mice have a normal distribution of TSH-producing pituitary cells, as well as normal T4 levels, therefore excluding abnormalities of thyroid function. The mechanism by which IL-6 causes a decrease in *in vivo* IGF-I levels remains to be clarified. Since circulating IGF-I is produced mainly by the liver (10), and IL-6 is involved in the regulation of gene transcription during inflammation (23, 35), it is tempting to hypothesize that overproduction of IL-6 may act negatively on liver IGF-I gene expression. On the other hand, since the great part of circulating IGF-I is carried in a ternary complex with IGF-binding protein-3 (IGFBP-3) and a non-IGF binding protein, termed acid labile subunit (ALS), and since the half-life of IGF-I is markedly prolonged by its association in this ternary complex from < 10 min to 12–15 h (10), it is also possible that a decrease in IGFBP-3 and/or in ALS may be responsible for the low circulating IGF-I levels.

Whatever the mechanism(s), our findings in mice strongly support the concept that IL-6 causes growth impairment in childhood chronic inflammatory diseases. Indeed, as previously mentioned, s-JRA is a chronic inflammatory disorder that associates markedly elevated circulating levels of IL-6 (12–17), stunted growth (2, 4, 5), and decreased IGF-I levels (7–9); in this study we found a negative correlation between circulating levels of IGF-I and IL-6, as well as between IGF-I levels and C-reactive protein concentrations. These findings, together with the results obtained in NSE/hIL-6 mice, strongly suggest that chronically elevated levels of IL-6 are responsible for the growth defect present in patients with s-JRA. Moreover, they appear to explain the observation that in patients with JRA treated with recombinant GH the height velocity during treatment was satisfactory in < 50% of the patients and was found to be inversely correlated with C-reactive protein concentrations (36). The possibility exists that, similarly to IL-6, other inflammatory cytokines, such as IL-1 or TNF- $\alpha$ , may affect IGF-I levels. However, in patients with s-JRA, we did not find any correlation of IGF-I levels with TNF- $\alpha$  or IL-1 $\beta$  levels.

Stunted growth associated with decreased levels of IGF-I has been reported in other diseases characterized by chronic inflammation or by chronic recurrent infections, including Crohn's disease and cystic fibrosis (37, 38). In Crohn's disease, growth retardation is related to inflammation (39). IL-6 is highly expressed in the inflamed mucosa and serum IL-6 levels are elevated and correlate with disease activity (40–42). In cystic fibrosis, growth retardation is significantly correlated with lung infections, but not with the degree of pancreatic insufficiency (43). Moreover, critically ill patients with sepsis, a condition with markedly elevated levels of IL-6, have substantially normal GH production, low IGF-I levels, and absent IGF-I response to exogenous GH, administered as anabolic agent (44, 45).

In conclusion, our study shows that in childhood chronic overproduction of IL-6 causes growth defect and that the growth defect appears to be mediated by a decrease in circulating IGF-I. These findings suggest that IL-6-mediated decrease

in IGF-I production may represent a generalized major mechanism by which chronic inflammation affects growth. This would imply that IGF-I and not GH represents the symptomatic treatment of choice for growth retardation in children with s-JRA, as well as in those with other chronic inflammatory diseases or with recurrent severe infections.

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