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Contrasting Skeletal Phenotypes in Mice with an Identical Mutation Targeted to Thyroid Hormone Receptor $\alpha 1$ or β

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Thyroid hormone (T₃) regulates bone turnover and mineralization in adults and is essential for skeletal development. Surprisingly, we identified a phenotype of skeletal thyrotoxicosis in T_3 receptor β^{PV} $(TR\beta^{PV})$ mice in which a targeted frameshift mutation in TR β results in resistance to thyroid hormone. To characterize mechanisms underlying thyroid hormone action in bone, we analyzed skeletal development in TR $\alpha 1^{PV}$ mice in which the same PV mutation was targeted to TR α 1. In contrast to $TR\beta^{PV}$ mice, $TR\alpha 1^{PV}$ mutants exhibited skeletal hypothyroidism with delayed endochondral and intramembranous ossification, severe postnatal growth retardation, diminished trabecular bone mineralization, reduced cortical bone deposition, and delayed closure of the skull sutures.

THE ACTIONS OF THYROID hormone (T₃) are mediated mainly by nuclear T₃ receptors (TRs), which act as ligand-inducible transcription factors (1). Several TR α and TR β isoforms are expressed in temporospatial patterns during development and in differing ratios in adult tissues (1–3). In the skeleton, TRs are expressed in growth plate chondrocytes, osteoblasts, and bone marrow stromal cells, and T₃ is essential for skeletal development, linear growth, and bone mineralization (4, 5). Childhood hypothyroidism causes growth arrest, delayed skeletal maturation, and epiphyseal dysgenesis, and T₄ replacement induces catch-up growth (6). Autosomal dominant resistance to thyroid hormone (RTH), which results from mutant TR β proteins, also impairs skeletal develop-

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Skeletal hypothyroidism in TR α 1^{PV} mutants was accompanied by impaired GH receptor and IGF-I receptor expression and signaling in the growth plate, whereas GH receptor and IGF-I receptor expression and signaling were increased in TR β^{PV} mice. These data indicate that GH receptor and IGF-I receptor are physiological targets for T₃ action in bone *in vivo*. The divergent phenotypes observed in TR α 1^{PV} and TR β^{PV} mice arise because the pituitary gland is a TR β -responsive tissue, whereas bone is TR α responsive. These studies provide a new understanding of the complex relationship between central and peripheral thyroid status. (*Molecular Endocrinology* 19: 3045–3059, 2005)

ment causing short stature and bone dysplasias (7, 8). Childhood thyrotoxicosis accelerates growth and advances bone age but induces short stature due to premature fusion of the growth plates. In severe cases, craniosynostosis can result from early closure of the skull sutures and may be associated with neurological deficits (9). Thus, euthyroidism is essential for normal bone development, and the skeleton is exquisitely sensitive to changes in thyroid status.

RTH is characterized by reduced responsiveness of target tissues to circulating T₄ and T₃. At the level of the hypothalamic-pituitary-thyroid axis this results in the classical biochemical features of RTH, in which negative feedback control of TSH production and secretion is impaired. The resulting inappropriately high levels of TSH drive increased T_4 and T_3 production, and a new equilibrium set point of high circulating concentrations of T₄ and T₃ with elevated levels of TSH is established. Mutations in the THRB gene cause RTH and result in the expression of mutant TR β proteins that are unable to respond normally to T₃, usually because of reduced binding affinity for T₃ or reduced transcriptional activation capabilities. The mutant $TR\beta$ proteins also interfere with the actions of normally expressed wild-type TR α and β and thus act as dominant-negative antagonists that disrupt transcription of

Abbreviations: ALSKO, Acid-labile subunit knockout; E17.5, embryonic d 17.5; FGFR, fibroblast growth factor receptor; GHR, GH receptor; IGF-1R, IGF-I receptor; LID, liver-specific IGF-I knockout; HZ, hypertrophic zone; P1, postnatal d 1; PZ, proliferative zone; RTH, resistance to thyroid hormone; RZ, reserve zone; STAT, signal transducer and activator of transcription; TR, T_3 receptor.

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T₃-regulated genes (10). The clinical syndrome of RTH is variable, resulting from both the direct effects of the mutant receptors and the consequences of elevated thyroid hormone concentrations. In addition, the differing ratios of TR α and TR β proteins that are expressed in individual tissues further complicate the syndrome. In tissues such as the heart, in which $TR\alpha$ predominates, the presence of tachycardia in RTH is likely to be due to the effects of elevated thyroid hormone levels acting via TR α . In contrast, in tissues such as the liver, in which TR β predominates, the presence of hypercholesterolemia in RTH may result from a combination of local tissue hypothyroidism and the direct dominant-negative actions of mutant TR β proteins in hepatocytes. Thus, the spectrum of clinical features in RTH is broad and can include reduced weight, cardiac disease, hypercholesterolemia, tachycardia, hearing loss, attention-deficit hyperactivity disorder, decreased IQ, and dyslexia (10-12). This complex spectrum reflects the presence of thyrotoxicosis in some T₃-target tissues but evidence of hypothyroidism in others.

A wide variety of bone phenotypes has been described in RTH, and this is probably because objective studies of the skeleton and growth are available in only a small minority of patients. Features include stippled epiphyses with scattered calcification in the growth plate, high bone turnover osteoporosis and fracture, reduced bone density, craniosynostosis, and various defects of facial bone and vertebral development (7). Growth retardation and short stature have been estimated to occur in 26% of patients with variably delayed bone age in up to 47% (7, 8), although bone age has also been shown to be advanced by more than 2 sps in two families, and lesser degrees of advancement have also been documented (7).

In a previous study, we characterized skeletal development in mutant mice with a PV mutation targeted to the TR β gene locus (13). The PV mutation was derived from a patient with severe RTH and consists of a C insertion at codon 448, which produces a frameshift of the carboxyl-terminal 14 amino acids of $TR\beta$ 1. The mutant $TR\beta^{PV}$ protein cannot bind T_3 , fails to transactivate T₃ target genes in vitro, and is a potent dominant-negative antagonist (14). $TR\beta^{PV}$ mice have very high levels of circulating T₄, T₃, and TSH (14), and we showed they display a phenotype of skeletal thyrotoxicosis (13). We also demonstrated that $TR\alpha 1$ mRNA is expressed at 12-fold higher concentrations than TR β 1 in bone, suggesting that the phenotype of skeletal hyperthyroidism in $TR\beta^{PV}$ mice results from increased T₃ levels acting via TR α 1 (13). To investigate this hypothesis and determine whether TR α 1 is functionally predominant in bone, we studied mice carrying the PV mutation targeted to TR α 1 (15). The mutant TR $\alpha 1^{PV}$ protein also acts as a potent dominant-negative antagonist that interferes with transcriptional activities of the wild-type TR α and TR β receptors (15). In keeping with other mice with dominant-negative RTH mutations in TR α 1 (11, 16), heterozygous TR α 1^{PV/+}

mice display only a modest degree of thyroid failure. There were small increases in TSH (1.7-fold) and T₃ (1.15-fold) levels in TR α 1^{PV/+} mice, but no change in circulating T₄ concentrations, and the homozygous mutation was lethal (15). Characterization of bone development in biochemically euthyroid TR α 1^{PV/+} mice, therefore, enabled us to investigate mechanisms of T₃ action in bone and determine whether TR α 1 is the major functional TR expressed in the skeleton.

RESULTS

$TR\alpha 1^{\text{PV}+}$ Mice Exhibit Delayed Ossification and Postnatal Growth Retardation

Analysis of bone lengths in TR $\alpha 1^{PV/+}$ mice revealed severe and persistent postnatal linear growth impairment (Fig. 1). No sexually dimorphic influences of the TR $\alpha 1^{PV}$ mutation on bone growth or development were observed. TR $\alpha 1^{PV/+}$ tibias were 15–25% shorter than tibias from wild-type littermates at all postnatal ages examined. In contrast, no difference was observed between embryonic d 17.5 (E17.5) and postnatal d 1 (P1) wild-type and TR $\alpha 1^{PV/+}$ mice. TR $\alpha 1^{PV/+}$





ulnas were also markedly shorter than wild-type (12–14% reduction), although the magnitude of growth impairment was less than in the tibia (Fig. 1). This degree of postnatal growth impairment in TR α 1^{PV/+} long bones was much more severe than documented in TR β ^{PV/+} and TR β ^{PV/PV} mice (13).

Analysis of skeletal preparations from E17.5 and P1 TR $\alpha 1^{PV/+}$ mice confirmed that bone lengths in mutant mice did not differ from wild type before birth, and the appearance of rib cages and vertebrae from wild-type and TR $\alpha 1^{PV/+}$ mice was similar (Fig. 2A). In these



Fig. 2. Skeletal Preparations from Wild-Type ($\alpha 1^{+/+}$) and TR $\alpha 1^{PV/+}$ Mice Stained with Alizarin Red (Bone) and Alcian Blue 8GX (Cartilage)

A, E17.5 mice, limbs ×10, rib cages ×7.5, and skulls ×11 magnification. *Arrows* indicate distal fore- and hindlimbs in TR α 1^{PV/+} mice and show reduced alizarin red staining in mutants compared with wild-type littermates. B, Neonatal mice, skulls ×11 and ×30 magnification. Anatomy of the skull sutures, bones, and fontanelles is shown in the *adjacent diagram*.

preparations, alizarin red stained ossified bone pink and alcian blue 8GX stained cartilage blue. Nevertheless, analysis of limbs in E17.5 mice revealed a small delay in endochondral ossification in the ulna and radius of the forelimb and in the tibia and fibula of the hindlimb that was evident in all TR $\alpha 1^{PV/+}$ mice examined (reduced alizarin red staining in these regions in $TR\alpha 1^{PV/+}$ mice [(n = 7) compared with wild-type (n = 6), arrowed in Fig. 2A]. Similar findings were present in neonatal mice (wild-type n = 3; TR $\alpha 1^{PV/+}$ n = 12; data not shown). These observations are in contrast with findings in TR $\beta^{PV/PV}$ mice, in which E17.5 and P1 skeletons displayed advanced endochondral ossification and were larger than wild-type littermates (13). Examination of the skull in E17.5 and neonatal mice revealed further differences between wild-type and $TR\alpha 1^{\mathsf{PV}/+}$ mice. There was no difference in anteriorposterior and biparietal skull dimensions, but the fontanelles in TR $\alpha 1^{PV/+}$ mice were larger and cranial sutures wider than in wild-type littermates [E17.5: 37.9 \pm 2.5 vs. 20.8 \pm 0.9 (P < 0.001); P1: 9.5 \pm 0.8 vs. 4.3 \pm 1.2 (P < 0.05), area of open fontanelles and sutures expressed as percentage of total skull area in TR $\alpha 1^{PV/+}$ vs. wild-type mice], indicating delayed fontanelle closure and suture fusion (Fig. 2). Intramembranous bone deposited in frontal and parietal bones of the $TR\alpha 1^{\mathsf{PV}/+}$ skull was also more porous and stained less intensely (Fig. 2B). These data demonstrate normal growth dimension, but markedly delayed intramembranous ossification of the skull in TR $\alpha 1^{PV/+}$ mice and contrast with findings in $TR\beta^{PV/PV}$ mice, in which advanced ossification of the skull with craniosynostosis was demonstrated (13).

Postnatal linear growth in TR α 1^{PV/+} and TR β ^{PV/PV} mice was examined in detail (Fig. 3). Tibias from $TR\alpha 1^{PV/+}$ mice were 15%, 17%, 20%, and 16% shorter than wild type at ages 2, 3, 4, and 7 wk, respectively, whereas tibias from $TR\beta^{PV/PV}$ mice were 11% shorter at 2 wk and only 2% shorter at 4 wk reflecting the accelerated growth spurt between these ages in TR $\beta^{PV/PV}$ mice (13). Growth impairment in $TR\alpha 1^{PV/+}$ mice was accompanied by reduced ossification of the secondary tibial epiphyses, a finding not seen in $TR\beta^{PV/PV}$ mice, but which persisted in $TR\alpha 1^{\mathsf{PV/+}}$ mice at 3 and 4 wk. Furthermore, hindlimb paws from $TR\alpha 1^{PV/+}$ mice at 3 and 7 wk revealed persistently impaired endochondral bone formation with delayed formation of secondary ossification centers in metatarsal bones at 2 wk and the presence of open metatarsal growth plates at 7 wk (Fig. 3). In contrast, epiphyseal ossification and metacarpal and metatarsal growth plate closure was advanced in 3 wk-old $TR\beta^{PV/PV}$ mice (13). These data indicate that postnatal linear growth impairment in TR $\alpha 1^{PV/+}$ mice was associated with delayed endochondral ossification, whereas in $TR\beta^{PV/PV}$ mice it resulted from advanced bone development.

Endochondral ossification in the proximal tibia was analyzed in histological studies in TR α 1^{PV/+} mice (Fig. 4). In wild-type mice the proximal tibia secondary os-



Fig. 3. Skeletal Preparations from Wild-Type ($\alpha 1^{+/+}$) and TR $\alpha 1^{PV/+}$ Mice Aged 2, 3, 4, and 7 wk and from Wild-Type ($\beta^{+/+}$), Heterozygote ($\beta^{PV/+}$), and Homozygous Mutant TR $\beta^{PV/PV}$ Mice Aged 2 and 4 wk

Tibias are shown at all ages and hindlimb paws are shown at ages 3 and 7 wk for $\alpha 1^{+/+}$ and $\alpha 1^{PV/+}$ mice. Preparations were stained with alizarin red (bone) and alcian blue (cartilage). Magnification ×6 in all cases. *Arrowheads* in panels showing tibias from 2-, 3-, and 4-wk mice indicate delayed formation of proximal and distal secondary ossification centers in $\alpha 1^{PV/+}$ mice. *Arrows* in panels showing hindlimb paws indicate delayed formation of secondary epiphyses (3 wk) and persistence of the growth plate (7 wk) in metatarsals of $\alpha 1^{PV/+}$ mice. The percentage differences in lengths of TR $\alpha 1^{PV/+}$ tibias compared with wild-type littermates, and TR $\beta^{PV/+}$ littermates, are shown.

sification center was already established with formation of bone trabeculae within the epiphysis at 2 wk. Between 2 and 7 wk there was progressive narrowing of the growth plate with increased epiphyseal trabecular bone deposition as endochondral ossification and bone maturation continued. In contrast, in TR $\alpha 1^{PV/+}$ mice endochondral ossification was markedly delayed. TR $\alpha 1^{PV/+}$ tibias were smaller in all dimensions, and development of the secondary ossification center was initiated at 3 wk, a time at which this process was well advanced in wild-type littermates. Formation of an organized growth plate, and subsequent growth plate



Fig. 4. Histological Sections of the Upper Tibia from Wild-Type (TR α 1^{+/+}) and TR α 1^{PV/+} Mice

A, Sections of the upper tibia from 2-, 3-, 4-, and 7-wk wild type (TR α 1^{+/+}) and TR α 1^{PV/+} mice (×50 magnification) stained with alcian blue (growth plate cartilage in blue) and van Gieson (bone osteoid in red). The proximal tibia secondary ossification center of the epiphysis (E) and the growth plate (GP) regions are shown along with immature epiphyses (IE). B, Undecalcified sections of the upper tibia and fibula from 2-wk wild type (TR α 1^{+/+}) and TR α 1^{PV/+} mice (×50 magnification) stained with von Kossa (calcified bone in *black*) and neutral *red* counterstain. The metaphysis (M) *below* the growth plate is shown and denotes the region in which calcified trabecular bone is stained *black*. The epiphysis in the wild type (TR α 1^{+/+}) contains calcified bone stained *black*, whereas no von Kossa staining is seen in this region in TR α 1^{PV/+} mice.

narrowing during the progression of endochondral ossification, was also markedly delayed. The histological features in 7-wk TR α 1^{PV/+} mice were similar to those in 3- to 4-wk wild-type mice, indicating that endochondral ossification was delayed by up to 4 wk in mutants (Fig. 4A). Delayed endochondral ossification in TR α 1^{PV/+} mice was associated with reduced deposition of calcified trabecular bone, as evidenced by reduced von Kossa staining of undecalcified sections of the tibia in 2-wk-old mutants (n = 3) compared with wild type (n = 3) (Fig. 4B). In particular, mineralization of trabecular bone in the secondary epiphysis in wildtype mice was already established by 2 wk of age, whereas in TR α 1^{PV/+} mice, staining in this region was absent. A smaller reduction in von Kossa staining was evident in trabecular bone in the region of the metaphysis in TR $\alpha 1^{PV/+}$ mice. In addition, cortical bone deposition in the tibial diaphysis was reduced by approximately 50% in 2-wk TR α 1^{PV/+} mice compared with wild-type littermates (Fig. 5). These data demonstrate that postnatal growth impairment in TR $\alpha 1^{PV/+}$ mice is associated with a 3- to 4-wk delay in bone formation, reduced trabecular bone mineralization, and impaired cortical bone deposition. The findings contrast with those in TR $\beta^{PV/PV}$ mice, in which advanced ossification and increased trabecular bone mineralization were evident (13).

To investigate mechanisms underlying delayed ossification in TR α 1^{PV/+} mice, measurements of specific regions in the growth plate were performed (Fig. 6). Histological studies enabled the reserve (RZ), proliferative (PZ), and hypertrophic (HZ) zones of growth plates to be identified (13, 17–19). In situ hybridization was performed to determine the expression of collagen II, a marker of proliferating chondrocytes (20), and allow measurement of growth plate dimensions (Fig. 6A). Between 2 and 4 wk there was progressive narrowing of the growth plate in wild-type mice that was due to the normal proportionate narrowing in each of the RZ, PZ, and HZ regions (Fig. 6B). The growth plate continued to narrow in wild-type mice between 4 and 7 wk, but at a slower rate than in younger animals reflecting growth plate maturation and its imminent quiescence as linear growth tails off toward adulthood (Fig. 6C). In TR $\alpha 1^{PV/+}$ mice, measurement of specific regions of the growth plate were not possible until animals reached 7 wk of age because, before that time, formation of the proximal tibial growth plate was incomplete (Fig. 4). At 7 wk the growth plate in



Fig. 5. Sections of the Midtibia Diaphysis from 2-wk-Old Wild-Type (TR α 1^{+/+}) and TR α 1^{PV/+} mice (×10 Magnification) Stained with van Gieson

Background shows nonspecific staining of bone marrow within the diaphysis cavity or skeletal muscle. The *arrow-heads* indicate the width of cortical bone present in wild-type TR α 1^{+/+} and mutant TR α 1^{PV/+} littermates.

 $TR\alpha 1^{PV/+}$ mice was significantly wider than in 7-wk wild-type littermates (P < 0.001) but did not differ in width when compared with growth plates from 4-wk wild-type mice. The finding that the width of the growth plate in 4-wk wild-type mice was similar to the width observed in 7-wk old $TR\alpha 1^{PV/+}$ mice suggested that endochondral ossification was delayed by about 3 wk in TR $\alpha 1^{PV/+}$ mice. Thus, comparisons of individual growth plate zones between 4-wk wild-type and 7-wk $TR\alpha 1^{PV/+}$ mice were made to investigate why ossification was delayed in TR $\alpha 1^{PV/+}$ mutants. These comparisons revealed that the PZ and HZ regions in $TR\alpha 1^{PV/+}$ mice were narrower than in wild-type (5%) and 10%, respectively; P < 0.05) but the RZ width was similar to wild type (Fig. 6B). Taken together, these data confirm that endochondral ossification in $TR\alpha 1^{PV/+}$ mice is delayed by approximately 3–4 wk and suggest this delay is due to impaired transition of immature RZ chondrocytes into the PZ, resulting in proportionally reduced numbers or dimensions of proliferating and hypertrophic chondrocytes. The data contrast with findings in TR $\beta^{PV/PV}$ mice, in which disproportionate and accelerated narrowing of the PZ and HZ regions accounted for premature growth plate quiescence by 4 wk of age (13).

TRα1^{PV/+} Mice Exhibit Skeletal Hypothyroidism

We previously identified that fibroblast growth factor receptor-1 (FGFR1) is a T₃-target gene in bone. Skeletal FGFR1 expression was reduced in TR α -null (TR $\alpha^{0/0}$) mice, which display a hypothyroid skeletal phenotype (19), but was increased in TR $\beta^{PV/PV}$ mice (13). In contrast, comparison of 3-wk-old wild-type mice with 7-wk-old TR α 1^{PV/+} mice (equivalent ages of growth plate maturation, Figs. 4 and 6) revealed that FGFR1 mRNA expression in both chondrocytes and osteoblasts was markedly reduced in TR α 1^{PV/+} mice (Fig. 7). These data demonstrate that TR α 1^{PV/+} mice display severe skeletal hypothyroidism.

GH and IGF-I Signaling in the Growth Plate Is Impaired in TR α 1^{PV/+} Mice, but Increased in TR β ^{PV} Mice

In view of the impaired transition of chondrocytes from RZ to PZ in TR α 1^{PV/+} mice, we investigated further by examining GH/IGF-I signaling in the growth plate. The GH/IGF-I pathway is initiated by GH, which activates the GH receptor (GHR) in growth plate chondrocytes. GH either acts directly on growth plate chondrocytes to regulate their proliferation and differentiation or stimulates local production of IGF-I, which subsequently acts in a paracrine manner to stimulate the IGF-I receptor (IGF-IR). IGF-I also exerts GH-independent actions on growth plate chondrocytes (21). GHR stimulation results in activation of a signaling cascade that involves signal transducer and activator of transcription (STAT)5 (22, 23), whereas stimulation of

Α









A, In situ hybridization for collagen II expression in proliferative zone of the upper tibia growth plate of a 3-wk WT mouse (×200 magnification). RZs, PZs, and HZs of the growth plate are indicated along with the regions of the secondary epiphysis (E) and primary spongiosum (PS). B, Graph showing relative widths of the RZ, PZ, and HZ regions and total growth plate heights (RZ+PZ+HZ) of 2-, 3-, and 4-wk WT mice compared with 7-wk TR α 1^{PV/+} mice. Mean growth plate height and zone width measurements (μ m) \pm SEM were obtained from two to three animals per group (three to four differing levels of section examined for each growth plate) by taking four separate measurements across each growth plate section examined. Data were analyzed by Student's t test to determine the differences in width of the growth plate zones and the total growth plate height between 7-wk TR α 1^{PV/+} mice and WT animals aged 2, 3, and 4 wk. C, *Graph* showing the decline in total growth plate height (μ m) \pm SEM with age in WT mice aged between 2 and 7 wk (n = 3 per group). NS, Nonsignificant (P = 0.63; P = 0.71).



Fig. 7. In Situ Hybridizations (\times 100 Magnification) for FGFR1 in Tibial Growth Plates from 4-wk Wild-Type (WT) and 7-wk TR α 1^{PV/+} Mice

The extent of the PZs and HZs are shown. Arrows indicate increased FGFR1 staining in osteoblasts lining trabecular bone surfaces within the secondary epiphysis in WT mice compared with TR α 1^{PV/+} mice.

IGF-IR results in activation of protein kinase B/Akt signaling (24, 25).

The GH/IGF-I pathway was investigated in growth plates from wild-type, $TR\alpha 1^{PV/+}$, $TR\beta^{PV/+}$, and TRBPV/PV mice by in situ hybridization and immunohistochemistry. In 4-wk wild-type mice, GHR was expressed at low levels only in prehypertrophic chondrocytes at the junction between the PZ and HZ. GHR expression was markedly increased in $TR\beta^{PV/PV}$ mice and extended throughout the PZ, whereas increased expression was also observed in $TR\beta^{PV/+}$ heterozygotes, but this was restricted to prehypertrophic chondrocytes. In contrast, GHR expression was absent from the growth plate in TR $\alpha 1^{PV/+}$ mice, although low levels of expression were evident in immature chondrocytes populating the incompletely formed growth plate from the region of the developing secondary epiphysis (Fig. 8). Low levels of IGF-I expression were also observed in these immature chondrocytes in TR $\alpha 1^{PV/+}$ mice, but not in the growth plate itself. In contrast, there were no differences in levels of IGF-I expression in $TR\beta^{PV/+}$ or $TR\beta^{PV/PV}$ mice compared with wild type, in which IGF-I mRNA was expressed in proliferating chondrocytes (Fig. 8). The patterns of expression of the IGF-IR were similar to those observed for expression of GHR. In wild-type animals IGF-IR was restricted to prehypertrophic chondrocytes. Expression was increased in the same region in $TR\beta^{PV/+}$ heterozygotes but was markedly increased throughout the growth plate in TR $\beta^{PV/PV}$ mice. In the TR $\alpha 1^{PV/+}$, IGF-IR expression was not detected in the growth plate but was present in immature chondrocytes located in the secondary epiphysis (Fig. 8).

To investigate whether the absence of GHR, IGF-I, and IGF-IR expression from growth plates in TR α 1^{PV/+} mice was because of the immaturity of the TR α 1^{PV/+} growth plate, we compared levels of expression in 2-, 3-, and 4-wk-old wild-type, TR α 1^{PV/+}, TR β ^{PV/+}, and TR β ^{PV/PV} mice (Fig. 9 and data not shown). Expression of all three mRNAs was absent from growth plates of TR α 1^{PV/+} mice at all ages but was present at low levels in immature chondrocytes in the region of the secondary epiphysis, consistent with findings in Fig. 8.



Fig. 8. In Situ Hybridizations (\times 200 Magnification) for GHR, IGF-I, and IGF-IR Expression in Tibial Growth Plates from 4-wk TR α 1^{PV/+}, Wild-Type (WT), TR β ^{PV/+} Heterozygote, and TR β ^{PV/PV} Mice

Note that $TR\beta^{+/+}$ wild-type mice are shown for simplicity; similar data were also obtained from $TR\alpha 1^{+/+}$ wild-type mice. The extent of the whole growth plate (GP) and PZs and HZs are shown. *Arrows* indicate positive staining in immature chondrocytes populating the incompletely formed immature growth plate in $TR\alpha 1^{PV/+}$ mice.

In wild-type mice, levels of GHR and IGF-IR decreased with age between 2 and 4 wk and became localized to prehypertrophic chondrocytes (Figs. 8 and 9 and data not shown). In contrast, in TR $\beta^{PV/+}$ and TR $\beta^{PV/PV}$ mice, expression of both GHR and IGF-IR remained persistently increased throughout the growth plate at all ages (Figs. 8 and 9 and data not shown). No changes in expression of IGF-I mRNA were observed in wild-type or mutant mice. These data indicate that altered patterns of expression of GHR and IGF-IR mRNAs in TR $\alpha 1^{PV/+}$, TR $\beta^{PV/+}$, and TR $\beta^{PV/PV}$ mice are not related to the maturity of the growth plate per se and suggest they result from altered skeletal T₃ signaling as a consequence of the TR $\alpha 1^{PV}$ or TR β^{PV} mutation.

To investigate whether changes in mRNA expression correlated with changes in functional activation of GHR and IGF-IR, we investigated the STAT5 and Akt downstream signaling pathways by immunohistochemistry. In TR α 1^{PV/+} mice, basal expression of STAT5 and Akt was no different than that of wild type, whereas concentrations of phosphorylated STAT5 and phosphorylated Akt were markedly reduced (Fig. 10). Thus, reduced levels of GHR and IGF-IR mRNAs in TR α 1^{PV/+} mice correlated with reduced activation of downstream signaling pathways. In TR β ^{PV/PV} mice, levels of basal STAT5 were reduced compared with wild type, whereas levels of basal Akt expression were unchanged. Concentrations of phos-

phorylated STAT5 and phosphorylated Akt, in contrast, were similar to wild type in TR $\beta^{PV/+}$ mice but were elevated in TR $\beta^{PV/PV}$ mice, although the increase in phosphorylated STAT5 was small (Fig. 11). Thus, increased expression of GHR and IGF-IR mRNAs in TR β^{PV} mice correlated with increased activation of downstream signaling pathways. Taken together, these data demonstrate that expression and activity of the local growth plate GH/IGF-I signaling pathway is markedly reduced in TR α 1^{PV} mice but increased in TR β^{PV} mice (Table 1), indicating that skeletal thyroid status is a key determinant of the sensitivity of growth plate chondrocytes to the local actions of GH and IGF-I.

DISCUSSION

We have demonstrated that TR α 1^{PV/+} mice exhibit a severe 25% reduction in postnatal linear growth, a 3-to 4-wk delay in endochondral ossification, diminished trabecular bone mineralization, reduced cortical bone deposition, and delayed intramembranous ossification. Reduced expression of the T₃-target gene FGFR1 indicates that skeletal hypothyroidism is responsible for this phenotype. The findings contrast with TR β^{PV} mice, in which skeletal thyrotoxicosis was documented by increased FGFR1 expression, accelerated



Fig. 9. *In Situ* Hybridizations (×200 Magnification) for IGF-IR Expression in Tibial Growth Plates from 2-, 3-, and 4-wk Wild-Type (TR $\beta^{+/+}$), Heterozygote TR $\beta^{PV/+}$, and TR $\beta^{PV/PV}$ Mice The extent of the PZs and HZs are shown.

early linear growth, increased trabecular bone mineralization, and advanced endochondral and intramembranous ossification that resulted in short stature and craniosynostosis (13). We previously identified that FGFR1 is a T₃-target gene in bone (19), and the major role of FGFR1 in skeletal development is to regulate intramembranous ossification of the skull (26). Activating mutations of FGFR1 cause Pfeiffer's craniosynostosis also occurs in severe childhood thyrotoxicosis (9). The presence of delayed closure of the skull sutures in TR $\alpha 1^{PV/+}$ mice, together with craniosynostosis in TR $\beta^{PV/PV}$ mice, suggests FGFR1 mediates important T₃ effects that regulate intramembranous ossification.

Nevertheless, prominent phenotypes in $TR\alpha 1^{PV/+}$ and TR β^{PV} mice involve abnormalities of endochondral bone formation. Thus, we investigated GH and IGF-I signaling in the growth plate, because they are major regulators of endochondral ossification and growth (21, 28, 29). The effects of GH were originally proposed to be mediated by liver-derived IGF-I (30), but this was challenged when IGF-I expression was identified in many tissues (21). A "dual effector theory" for GH action was proposed (31) and extrapolated to the growth plate (32). In this model GH initiates differentiation of PZ chondrocytes directly and is proposed to induce local IGF-I production in proliferating chondrocytes. Local IGF-I then acts in an autocrine/paracrine manner to stimulate clonal expansion and chondrocyte proliferation, resulting in longitudinal growth. However, evidence from IGF-I knockout (IGF- $I^{-/-}$),

liver-specific IGF-I knockout (LID), acid-labile subunit knockout (ALSKO) (acid-labile subunit forms a ternary complex with IGF-I and IGF-I binding protein-3 to stabilize serum IGF-I and facilitate its endocrine actions), and LID+ALSKO double-knockout mice have demonstrated that a threshold concentration of circulating IGF-I is also necessary for bone growth. Nevertheless, tissue IGF-I also plays an essential role because IGF-I^{-/-} mice are much more growth retarded than LID+ALSKO double-knockout mice (33-37). Data from GHR knockout (GHR^{-/-}), IGF-I^{-/-}, and GHR^{-/-}IGF-I^{-/-} double mutants indicate that GH and IGF-I act on the growth plate by both independent and overlapping pathways, with IGF-I being the major determinant of embryonic and postnatal growth, and its actions being modulated by GH in the postnatal period (29). It has further been suggested that, because only 17% of somatic growth can be attributed to processes that do not require an intact GH/IGF-I axis, GH and IGF-I pathways in the growth plate act as a point of convergence and participate in the actions of most growth-promoting molecules (29).

This concept is supported by studies showing that IGF-I is stimulated by T_3 in osteoblastic cells (38, 39) and IGF-IR is T_3 responsive in chondrocyte cultures (40). A recent study also showed that T_3 treatment of hypophysectomized rats resulted in increased GHR expression in the growth plate (41), although a previous study showed that GHR expression in the growth plate was independent of thyroid status (42). In addi-



Fig. 10. Immunohistochemistry (×200 Magnification) of STAT5 (*Upper Panels*), pSTAT5 (*Second Row*), Akt (*Third Row*) and pAkt (*Fourth Row*) ProteinExpression in Tibial Growth Plates from Wild-Type (TR α 1^{+/+}) and TR α 1^{PV/+} Mice

The *right-hand column* labeled "Controls" shows parallel experiments in which primary antibody was omitted from the immunohistochemistry protocol and which show that staining of STAT and Akt proteins occurred only in the presence specific primary antibody. pSTAT5, Phosphorylated STAT5; pAkt, phosphorylated AKT.

tion to effects on local growth plate GH/IGF-I signaling, T_4 and T_3 influence pituitary GH secretion (43). Abnormalities of the GH/IGF-I axis have been documented in various TR knockout mice: TR $\alpha 1^{-/-\beta-/-}$ mice (which lack TR α 1 and TR β) have GH- and mild IGF-I deficiency (44), and GH replacement restores their growth but does not improve defective ossification (45): $TR\alpha^{0/0\beta-1/-}$ mice (lacking all products of the *Thra* and Thrb genes) have GH deficiency (17); $TR\alpha^{0/0}$ mice have normal GH production (17); $TR\beta^{-/-}$ mice have mildly reduced GH production (46); and TR $\alpha 2^{-/-}$ mice (which lack TR α 2 but overexpress TR α 1) have normal GH levels but are IGF-I deficient (47). We previously showed that $TR\alpha 1^{PV/+}$ mice have normal pituitary GH production (15), whereas GH expression is reduced by 80% and circulating IGF-I is reduced by 40% in TR $\beta^{PV/PV}$ mutants (14, 48). These data from various TR mutant mice indicate that T_3 , acting mainly via TR β , regulates systemic GH/ IGF-I signaling pathways in vivo. Nevertheless, the presence of delayed endochondral ossification in $TR\alpha 1^{PV}$ mice despite normal levels of GH, and the presence of accelerated ossification in $TR\beta^{PV/PV}$ mice in the face of low levels of GH and IGF-I, is discordant with the known actions of GH/IGF-I in the growth plate. These findings strongly suggest that the skeletal consequences of the PV mutation result from dysregulated local GH/IGF-I signaling in the growth plate.

Data in Figs. 8-11 support this by clearly showing that GHR and IGF-IR expression and signaling are reduced in TR $\alpha 1^{PV}$ mice (skeletal hypothyroidism) but increased in TR β^{PV} mice (skeletal thyrotoxicosis). Nevertheless, the increase in GHR expression in $\text{TR}\beta^{\text{PV/PV}}$ growth plates (Fig. 8) was accompanied by a disproportionately small rise in activated STAT5 (Fig. 11). This finding reflects the impaired GH production observed in these mice (14) and demonstrates that the net effect of the $\text{TR}\beta^{\text{PV}}$ mutation results from systemic and local consequences on GH action. In contrast, IGF-I expression was unchanged in $TR\beta^{PV/+}$ and $TR\beta^{PV/PV}$ mice compared with wild type and was undetectable in TR $\alpha 1^{PV}$ growth plates, suggesting that TR α and/or GHR activity are necessary for IGF-I expression but indicating that growth plate IGF-I expression is not responsive to increased T_3 or GH action.



Fig. 11. Immunohistochemistry (×200 Magnification) of STAT5 (*Upper Panels*), pSTAT5 (*Second Row*), Akt (*Third Row*), and pAkt (*Fourth Row*) ProteinExpression in Tibial Growth Plates from Wild-Type (TRβ^{+/+}), TRβ^{PV/+}, and TRβ^{PV/PV} Mice The *right-hand column* labeled "controls" shows parallel experiments in which primary antibody was omitted from the immunohistochemistry protocol and which show that staining of STAT and Akt proteins occurred only in the presence of specific

primary antibody. pSTAT5, Phosphorylated STAT5; pAkt, phosphorylated AKT.

Nevertheless, increased activation of Akt was observed in TR $\beta^{PV/PV}$ mice and was independent of changes in IGF-I expression, instead correlating with increased IGF-IR expression. These findings suggest that levels of IGF-IR, rather than IGF-I ligand, are limiting in the growth plate. An alternative possibility is that an unidentified T₃-stimulated, IGF-I independent pathway could increase Akt activation in TR $\beta^{PV/PV}$ mice. Taken together, these data indicate that local GH/IGF-I actions mediate important effects of T₃ on endochondral ossification.

Nevertheless, in TR α 1^{PV/+} mice with skeletal hypothyroidism and reduced GHR and IGF-IR activity, growth plates were observed to be wider than in wildtype mice (Fig. 4), whereas in TR $\beta^{PV/PV}$ mice, with skeletal thyrotoxicosis and increased GHR and IGF-IR signaling, growth plates were narrower (13). In contrast, growth plates were observed to be narrower in GHR^{-/-} and IGF-I^{-/-} mice compared with wild type

(29, 36, 49), suggesting that T₃ exerts important effects on linear growth that are independent of GH and IGF-I. Indeed, TR α and TR β are expressed in growth plate chondrocytes (50-53). T₃ inhibits clonal expansion and proliferation but promotes hypertrophic differentiation of primary chondrocytes in suspension culture (53), and additional studies have shown T_3 regulates the spatial organization of chondrocyte columns and is required for terminal hypertrophic differentiation (54). In contrast, IGF-I stimulates chondrocyte proliferation and differentiation (28). Furthermore, growth retardation in hypothyroidism results from disrupted growth plate architecture, impaired vascular invasion of the growth plate, and inhibition of hypertrophic chondrocyte differentiation (18, 55). Again, differences are apparent as growth retardation in GH and IGF-I deficiency results from a combination of impaired chondrocyte proliferation and a reduction in the linear dimension of terminal hypertrophic chondro-

	Pituitary GH Production	Growth Plate				
		GHR mRNA	GH Signaling (pSTAT5)	IGF-I mRNA	IGF-IR mRNA	IGF-I Signaling (pAkt)
α1PV	Normal	Reduced	Reduced	Reduced	Reduced	Reduced
βPV	Reduced	Increased	Increased	Normal	Increased	Increased

cytes (28, 29, 36, 49, 54). Together, these considerations indicate that regulation of growth and endochondral ossification by T_3 involves both GH/IGF-Iindependent and GH/IGF-I-dependent pathways.

In these studies, we showed that $TR\alpha 1^{PV/+}$ mice display skeletal hypothyroidism despite the presence of biochemical euthyroidism. In contrast, $TR\beta^{PV}$ mice have severe RTH but a phenotype of skeletal thyrotoxicosis (13). This paradox results from differing effects of the PV mutations at the level of the hypothalamic-pituitary-thyroid axis and in bone (Fig. 12). The hypothalamus and pituitary predominantly express TR β , and mutation or deletion of TR β results in impaired feedback regulation of TSH and the syndrome of RTH with thyrotoxic levels of T₄ and T₃ and elevated TSH concentrations (14, 44, 46, 56-58). In this situation, the pituitary displays tissue hypothyroidism. In contrast, mutation or deletion of TRa does not interfere significantly with feedback regulation of TSH, and minor changes in circulating T_4 and T_3 levels result from impaired hormone production in the thyroid gland (11, 15-17, 58, 59). In this situation the pituitary functions normally and systemic T₄ and T₃ levels lie within or close to the normal range. Together with data from other mutant mice (reviewed in Refs. 12, 60, and 61), these considerations establish that $TR\beta$ is the physiological mediator of negative feedback control of TSH secretion. In contrast, our previous studies suggest that TR α is the major functional TR in bone (13,

17, 19). In the current studies, demonstration of skeletal hypothyroidism and impaired ossification in TR α 1^{PV/+} mice establishes that TR α acts directly in bone as a physiological regulator of skeletal development. In this context, it is apparent that the skeletal consequences of disrupted TR β function in TR β^{PV} mice result from impaired inhibition of TSH and the resulting elevated T₄ and T₃ concentrations, which act via TR α in bone to induce skeletal thyrotoxicosis (13, 62). In contrast, the skeletal hypothyroidism in TR α 1^{PV/+} mice results from locally impaired TR α function in bone.

Our analysis of TR $\alpha 1^{PV/+}$ and TR β^{PV} mice has provided a new understanding of the complex relationship between central pituitary thyroid status and peripheral skeletal thyroid status that arises because the pituitary gland is a TR β target tissue, whereas bone is a TR α target organ. The model in Fig. 12 can be extended to understand the relationship between central and peripheral thyroid status in any T₃-target tissue, depending on whether the peripheral tissue in question is TR α or TR β responsive. Thus, in the heart, a TR α -responsive organ, features of thyrotoxicosis are seen in mice with TR β mutation (63), whereas features of hypothyroidism are seen in TR α mutants (59). In contrast, in the liver, a TR β target tissue, a hypothyroid phenotype of impaired cholesterol clearance is seen in TR β mutant mice but not in TR α mutants (64).



Fig. 12. Relationship between Pituitary and Skeletal Thyroid Status Revealed by Analysis of $TR\alpha 1^{PV/+}$ and $TR\beta^{PV/PV}$ Mice Bone is a $TR\alpha$ -responsive tissue, whereas pituitary is $TR\beta$ responsive. The $TR\alpha 1^{PV}$ mutation does not affect pituitary T_3 responses because mutant $TR\alpha 1^{PV}$ concentrations are too low to interfere with $TR\beta$. In bone, however, T_3 responses are severely impaired because of high concentrations of dominant-negative $TR\alpha 1^{PV}$, resulting in skeletal hypothyroidism. In contrast, the $TR\beta^{PV}$ mutation disrupts pituitary T_3 responses and causes RTH, resulting in elevated circulating thyroid hormone concentrations and reduced GH production. In bone, mutant $TR\beta^{PV}$ concentrations are too low to interfere with $TR\alpha 1$, and elevated T_4 and T_3 concentrations hyperstimulate $TR\alpha 1$, resulting in skeletal thyrotoxicosis. TRE, Thyroid response element.

MATERIALS AND METHODS

$TR\alpha 1^{PV}$ and $TR\beta^{PV}$ Mutant Mice

Animal studies were conducted in strict accordance with the National Institutes of Health Guide for Care and Use of Laboratory Animals and were approved by the National Cancer Institute Animal Care and Use Committee. Wild-type, heterozygous (TR α 1^{PV/+} and TR β ^{PV/+}), and homozygous $(TR\beta^{PV/PV})$ mutant mice were bred and genotyped as described elsewhere (14, 15). Both $TR\alpha^{1PV}$ and $TR\beta^{PV}$ strains were generated from genomic clones isolated from a 129Sv mouse genomic library and transfected into TC-1 embryonic stem cells. Both mutant strains have a mixed C57BL/6J and NIH Black Swiss genetic background. Initial studies have revealed that $TR\alpha 1^{PV/+}$ mice exhibit growth retardation with only minor alterations in circulating thyroid hormones (15). In contrast, gross RTH is present in homozygous $TR\beta^P$ mice, with milder thyroid dysfunction in heterozygous $TR\beta^{PV/+}$ mutants (14). Detailed analysis of bone development in ${\rm TR}\beta^{\rm PV}$ mice has revealed accelerated growth, advanced bone age, and short stature resulting from skeletal thyrotoxicosis (13).

Skeletal Preparations

E17.5, P1, P14, P21, P28, and P49 male and female littermate mice were obtained. E17.5 and neonatal mice and limbs from P14, P21, P28, and P49 animals were fixed in 95% ethanol before staining with alizarin red and alcian blue 8GX as described previously (13). Skeletal preparations were photographed using a Leica MZ 75 binocular microscope (Leica AG, Heerbrugg, Switzerland), Leica KL 1500 LCD light source, Leica DFC 320 digital camera, Leica IM50 Digital Image Manager, and Leica Twain Module DFC 320 image acquisition software. Bone lengths from wild-type and TR $\alpha 1^{PV/+}$ male and female littermates were determined digitally after linear calibration of pixel size using the image acquisition software. Skull dimensions and open fontanelle and suture areas were calculated using Image J v1.33u software (http://rsb.info.nih.gov/ij/). The assessment of ossification stage in E17.5 and neonatal mice, as determined by the amount of alizarin red staining relative to alcian blue, was more subjective (Fig. 2). In these studies differences that were observed in all mutant mice examined compared with wild-type littermates were considered to be indicative of a difference in the degree of ossification.

Histology

Limbs were fixed for 48–72 h in 10% neutral buffered formalin followed by decalcification in 10% formic acid and 10% neutral buffered formalin at 20 C. E17.5 and P1 limbs were decalcified for 24 h; P14, P21, and P28 limbs were decalcified for 5 d and P49 limbs were decalcified for 7 d. Paraffinembedded 3- μ m sections were taken from anatomically oriented bones (three to five parallel levels per bone depending on the age of the animal; 20 sections per level) and stained with hematoxylin and eosin (Pioneer Research Chemicals, Colchester, UK) or van Gieson and alcian blue 8GX, as described elsewhere (13, 18). Some limbs from E17.5 and P14 mice were also fixed for 48–72 h in 10% neutral buffered formalin and frozen in paraffin without prior decalcification for determination of mineralization by von Kossa staining of 3- μ m cryosections with neutral red counterstain (13).

In Situ Hybridization and Analysis of Growth Plate Dimensions

mRNA expression was analyzed in growth plate sections from P14, P21, P28, and P49 mice using collagen II, collagen

X, FGFR1, IGF-I, IGF-IR, and GHR cRNA probes. A bacterial neomycin resistance gene cRNA probe (Boehringer Mannheim. Lewes. Sussex. UK) was used as a negative control for all hybridizations, and collagen II (nucleotides 2982-3689; GenBank accession no. L48440) and X (nucleotides 418-858; GenBank accession no. AJ31848) probes were used to identify proliferative and hypertrophic zones in growth plate sections, as described in previous studies in which we optimized in situ hybridization methods (13, 18, 19). A rat FGFR1 (nucleotides 104-603; GenBank accession no. S54008) partial cDNA was isolated by RT-PCR as described previously (18, 19) from osteoblastic ROS 17/2.8 cells (65). The rat IGF-I partial cDNA (nucleotides 61-314; GenBank accession no. D00698) was a gift from Dr. Cécile Kedzia (Institut National de la Santé et de la Recherche Médicale, Paris, France). Mouse IGF-1R (nucleotides 1063-1690; GenBank accession no. XM_133508) and GHR (nucleotides 470-711; GenBank accession no. NM_010284) partial cDNAs were isolated by RT-PCR as described elsewhere (18, 19) from chondrogenic ATDC5 cells (66) with the following primers: IGF-1R, forward 5'-GAAGACCACCATCAACAAT-3', reverse 5'-GAAGGA-CAAGGAGACCAAG-3'; GHR, forward 5'-GACCCCAG-GATCTATTCAGC-3', reverse 5'-CAGGTTGCACTATTTCGT-CAAC-3'. PCR products were subcloned into pGEM-T (Promega, Southampton, Hampshire, UK) and sequenced. FGFR1, IGF-I, IGF-1R, and GHR constructs were linearized with Spel, BamHI, Drall, and Spel, and digoxigenin-labeled cRNA probes were synthesized using T7, T3, T7, and T7 RNA polymerases, respectively (Boehringer Mannheim). In situ hybridizations using alkaline phosphatase-labeled probes were performed on 3-µm deparaffinized sections as described elsewhere (13, 18, 19). Studies were performed on at least three mice per genotype in duplicate, and repeat experiments

were performed on three separate occasions. Measurements at four separate positions across the width of growth plates were obtained, using a Leica DM LB2 microscope, Leica DFC 320 digital camera, Leica IM50 Digital Image Manager, and Leica Twain Module DFC 320 image acquisition software, to calculate mean values for the heights of the RZ, PZ, HZ, and total growth plate in sections from wild-type and TR α 1^{PV/+} mice. Results from adjacent levels of sectioning were compared to ensure consistency of the data. Cortical bone width measurements were performed at four separate positions in the midshaft of the tibia and adjacent levels of sectioning were compared. All studies are performed with the observer blinded to the genotype. For histology and histomorphometry analyses, at least three animals per genotype were examined.

Immunohistochemistry

Activation of IGF-IR and GHR downstream signaling was examined by immunohistochemical analysis of protein kinase B (Akt) and signal transducer and activator of transcription-5 (STAT5) expression in wild-type, $TR\alpha 1^{PV/+}$, $TR\beta^{PV/+}$, and $TR\beta^{PV/PV}$ growth plates. Sections were deparaffinized and rehydrated in ethanol and PBS. Sodium citrate antigen retrieval was performed for 6 min in a microwave oven on medium setting. Endogenous peroxidase activity was quenched with 1% H₂O₂ in methanol for 15 min at room temperature. Sections were then blocked with 5% fetal calf serum (Sigma Chemical Co., St. Louis, MO) in PBS with 0.5% Tween 20 for 1 h at room temperature, before addition of primary antibody and incubation overnight at 4 C. Polyclonal antibodies used to detect expression of Akt (Santa Cruz Biotechnology. Inc., Santa Cruz, CA), phosphorylated Akt (Cell Signaling Technology, Inc., Beverly, MA), STAT5 (Santa Cruz), and phosphorylated STAT5 (Santa Cruz) were diluted 1:175, 1:50, 1:200, and 1:140, respectively. Sections were subsequently incubated with peroxidase-conjugated secondary antibody (Bio-Rad Laboratories, Inc., Hercules, CA) diluted 1:2000 to 1:1800 for 30 min at room temperature. Peroxidase activity was detected using 3,3'-diaminobenzidine containing 0.02% H_2O_2 (Sigma). Negative controls lacking primary antibody were performed in parallel in all experiments, as described elsewhere (53, 67). Studies were performed on at least three mice per genotype in duplicate, and repeat experiments were performed on three separate occasions.

Statistical Analysis

Data were expressed as mean \pm sEM. Differences between groups were examined for statistical significance using Student's *t* test, in which *P* values <0.05 were considered significant.

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