

REVIEW

Role and Mechanisms of Actions of Thyroid Hormone on the Skeletal Development

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The importance of the thyroid hormone axis in the regulation of skeletal growth and maintenance has been well established from clinical studies involving patients with mutations in proteins that regulate synthesis and/or actions of thyroid hormone. Data from genetic mouse models involving disruption and overexpression of components of the thyroid hormone axis also provide direct support for a key role for thyroid hormone in the regulation of bone metabolism. Thyroid hormone regulates proliferation and/or differentiated actions of multiple cell types in bone including chondrocytes, osteoblasts and osteoclasts. Thyroid hormone effects on the target cells are mediated via ligand-inducible nuclear receptors/transcription factors, thyroid hormone receptor (TR) α and β , of which TR α seems to be critically important in regulating bone cell functions. In terms of mechanisms for thyroid hormone action, studies suggest that thyroid hormone regulates a number of key growth factor signaling pathways including insulin-like growth factor-I, parathyroid hormone related protein, fibroblast growth factor, Indian hedgehog and Wnt to influence skeletal growth. In this review we describe findings from various genetic mouse models and clinical mutations of thyroid hormone signaling related mutations in humans that pertain to the role and mechanism of action of thyroid hormone in the regulation of skeletal growth and maintenance.

Keywords: thyroid hormone; bone; cartilage; growth factors; bone cells

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Introduction

Thyroid hormone (TH) plays an important role in normal endochondral ossification and is essential for skeletal development, linear growth, maintenance of bone mass, and efficient fracture healing (1). Juvenile hypothyroidism causes growth arrest with delayed bone formation and mineralization, and T4 replacement induces rapid catch-up growth (2). By contrast, childhood thyrotoxicosis accelerates bone formation with premature closure of the growth plates and skull sutures, leading to short stature and craniosynostosis (3). Although there is

considerable evidence regarding the importance of TH in skeletal development, the molecular mechanisms of TH action in bone are poorly understood. In this chapter, we discuss regulation and mechanisms of action of TH during skeletal development with particular emphasis on areas in which recent advances have been made.

Physiology of TH: Regulation, metabolism and TH receptor

Regulation

Systemic TH levels are maintained by the classical negative feedback loop involving the hypothalamus-pituitary-thyroid (HPT) axis (Figure 1). Thyrotropin releasing hormone (TRH) is synthesized in the paraventricular nucleus (PVN) of the hypothalamus and stimulates synthesis and secretion of thyroid stimulating hormone (TSH) from

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thyrotroph cells in the anterior pituitary gland. TSH subsequently acts via the TSH receptor (TSHR) on thyroid follicular cells to stimulate synthesis and release of 3,5,3',5'-L-tetraiodothyronine (thyroxine, T4) and 3,5,3'-L-triiodothyronine (T3). The circulating T4 and T3 are predominantly bound to carrier proteins including thyroxine binding globulin, transthyretin (previously known as thyroxine binding pre-albumin) and albumin, with only approximately 0.2% of the total T3 and 0.02% of the total T4 available as free unbound hormones (fT3, fT4) in plasma.

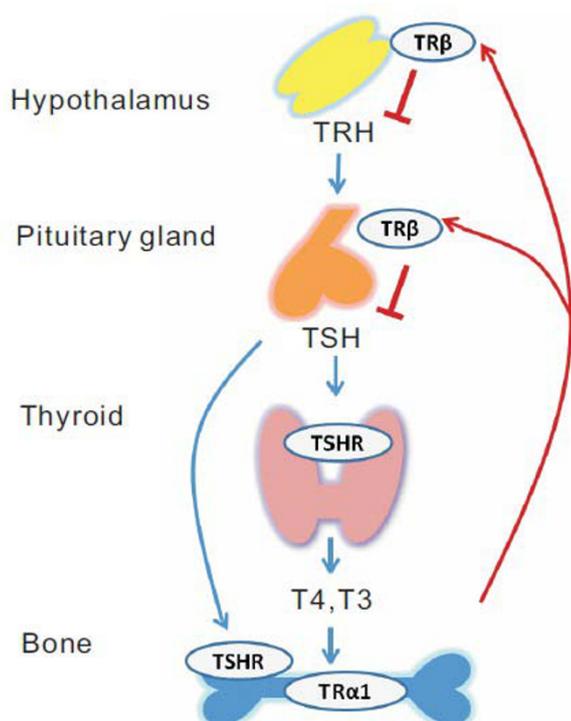


Figure 1 TRH-TSH-T3 feedback loop. The hypothalamic neurons secrete thyrotropin releasing hormone (TRH) which is carried down to the adenohypophysis of the pituitary by the hypothalamic portal vein where it releases thyroid stimulating hormone (TSH). The released TSH reaches thyroid glands via blood stream to bind to TSH receptor (TSHR) to stimulate production and release of thyroxine (T4) and T3. T3 exerts its actions on bone mainly by binding to TR α . TSH can also act directly on bone cells by binding to TSHR. Increased levels of T3 can act by negative feedback loop via TR β to inhibit release of TRH and TSH, thereby preventing hyperparathyroidism.

TH is known to act via the nuclear TH receptor β (TR β) in the hypothalamus and pituitary to inhibit TRH and TSH production and secretion (4-5) and thus complete a negative feedback loop that maintains systemic thyroid status within a normal reference range. This negative feedback loop maintains a physiological inverse relationship between TSH and circulating T3 and T4 levels that

defines the HPT axis set-point (6-7).

Metabolism

The predominant circulating TH is the pro-hormone T4, which can be converted to the biologically more potent hormone, T3. TH metabolism is mediated by three iodothyronine deiodinases. The type 1 and type 2 enzymes (D1 and D2) convert T4 to T3 by catalyzing removal of a 5'-iodine atom. By contrast, the type 3 enzyme (D3) irreversibly removes a 5-iodine atom from either T4 or T3 to generate the inactive metabolites 3,3',5'-L-triiodothyronine (reverse T3, rT3) and 3,3'-diiodothyronine (T2), respectively (8-9). D1 is not expressed in skeletal cells (10-11), indicating D1 does not influence T3 action on bone directly. D2 is restricted to mature primary osteoblasts but is undetectable in chondrocytes and osteoclasts (12).

The cellular influx as well as efflux of iodothyronines is known to be mediated by several specific membrane transporter proteins including the monocarboxylate transporters 8 and 10 (MCT8 and MCT10), sodium-dependent organic anion co-transporting polypeptide 1 (OATP1), the sodium taurocholate co-transporting polypeptide (NTCP) and the L-type amino acid transporter 1 (LAT1) and LAT2 (13-15). A study by Capelo *et al* revealed that MCT8, LAT1 and LAT2 are expressed in the skeletal tissues of mice as well as in osteoblastic MC3T3-E1 cells (16). Thus, the intra-cellular levels of the active hormone, T3, and its availability to nuclear TH receptors (TRs) are determined by the relative activities of D2 and D3 as well as expression levels of TH transport proteins.

TH receptor/ TH action

The major action of TH is exerted through nuclear TH receptors (TRs), which are ligand-inducible transcription factors. Based on chromosomal localization and amino acid homology, two classes of TRs, α and β , have been identified. Due to differential splicing of these two genes, multiple TRs are generated as α 1, α 2, α 3, β 1, β 2, and β 3, as well as three truncated forms, $\Delta\alpha$ 1, $\Delta\alpha$ 2, $\Delta\beta$ (17-18). The α 2 and α 3 isoforms and all of the truncated receptors are non-T3 binding proteins that function as antagonists of TH signaling (18-20). TR α 1 and TR β 1 are expressed in virtually all tissues, but their abundance and roles differ, depending on the developmental stage of the organism and on the particular tissue type (21). TR α 1 is more abundantly expressed in heart, brain, and bone, while TR β 1 is more highly expressed in liver and pituitary (22). By contrast, expression of TR β 2 is restricted to the hypothalamus and pituitary where it mediates inhibition of TRH and TSH expression and the cochlea and retina where it regulates sensory organ development (23-24)

and TR β 3 is expressed in kidney, liver, and lung (25). Thus, TH action in target tissues is determined in part by the types and abundance of TH receptors present.

In the nucleus, TRs form homodimers with another TR or heterodimers with retinoid X receptors (RXR) and bind to specific TH response element sequences (TREs) located in promoter regions of T3-target genes and regulate their expression in a ligand-dependent manner. Unliganded TRs bind TREs in T3 target genes and mediate transcriptional repression. Co-repressor proteins such as nuclear receptor corepressor protein/silencing mediator of retinoid and TH receptors are recruited to the RXR-TR heterodimer in the absence of T3 and inhibit target gene expression. T3 binding displaces the co-repressor, allowing co-activator proteins such as CBP/p300, pCAF, and SRC-1 to interact with the RXR-TR heterodimer and activate gene transcription in a hormone-dependent manner (26-28).

Besides the genomic actions of T3, nongenomic mechanism of TH analogues are increasingly recognized to have downstream consequences at the level of specific gene transcription (26, 29). The nongenomic mechanisms of TH are known to be initiated at the plasma membrane, in the cytoplasm or in the intracellular organelles, such as mitochondria. At the membrane level, TH may interact with integrin α V/ β 3 to activate ERK1/2 which culminates in regulation of ion transport systems or cell proliferation (30). The relative contribution of nongenomic mechanisms in mediating TH effects on skeletal development is yet to be determined.

Skeletal development

The skeleton in different parts of the body develops through two distinct processes, intramembranous ossification and endochondral ossification. Intramembranous ossification, which occurs in the flat bones of the skull, involves direct differentiation of embryonic mesenchymal cells into bone-forming osteoblasts without an intermediate cartilage model (31). By contrast, endochondral ossification, which occurs in the remainder of the skeleton, involves the replacement of a cartilage model by bone tissue. Mesenchymal precursor cells condense and differentiate into chondrocytes, which secrete matrix proteins to form a cartilage template. The model expands through chondrocyte proliferation. Ossification of the cartilage model is preceded by hypertrophy of the chondrocytes in the prospective mid-shaft of the bone. Subsequently, blood vessels, osteoclasts (cartilage- and bone-resorbing cells), as well as bone marrow and osteoblast precursors then invade the model from the bone collar and proceed to form the primary ossifica-

tion center. The primary center expands towards the ends of the cartilage model, as the osteoclasts remove cartilage extracellular matrix (ECM) and osteoblasts deposit bone on cartilage remnants. In long bones, secondary ossification centers form within cartilage at the ends of long bones and remain separated from the primary ossification center by the epiphyseal growth plates. Skeletal maturity occurs when the expanding primary ossification center meets the secondary ossification center, thus obliterating the growth plate. The ordered process of growth plate chondrocyte proliferation, hypertrophic differentiation, apoptosis and subsequent new bone formation mediates linear growth until adulthood (32).

The process of endochondral ossification and the rate of linear growth are tightly regulated by multiple systemic hormones (including THs, growth hormone (GH), glucocorticoids and sex steroids) and various cytokines and growth factors (including insulin-like growth factor 1 (IGF-1), parathyroid hormone-related peptide (PTHrP), Indian hedgehog (Ihh), bone morphogenetic protein (BMP)s, fibroblast growth factor (FGF)s and vascular endothelial growth factor (VEGF)s that act in a paracrine and autocrine manner (33). The interaction between systemic hormones and local factors play a critical role in regulating linear growth, bone mass accumulation and mineralization processes until peak bone mass (34) is achieved in early adulthood. Throughout adult life there is a gradual loss of bone mass, which in women is accelerated at the menopause. Excessive TH as in the case of hyperthyroidism in adults is known to induce increased bone turnover and fracture incidence (35-36). While TH is known to be involved in regulating development of peak bone mass during early childhood and in the maintenance of bone mass in adults, this review will focus on the mechanism of TH action in regulating skeletal development.

Effects of TH on skeletal development

A number of genetic mouse models (37-38) have been generated to date to elucidate the role of the TH axis on growth and development of a number of tissues including bone. The skeletal phenotypes of these mutant mouse models will be described below.

Skeletal phenotype of TH signaling related mutant mice (Table 1)

Mouse mutants with altered TSH or TH levels

Pax8^{-/-} mice lack the thyroid specific transcription factor Pax8 required for thyroid follicular cell formation (39-40) and *hyt/hyt* mice have a loss-of-function mutation in the

Table 1 Skeletal phenotype of TH signaling related mutant mice

KO	Deleted Protein	Skeletal Phenotype	TH-pituitary axis	Reference
Pax8	Lack of essential transcription factor Pax8 for thyroid follicular cell development	Impaired linear growth, delayed endochondral ossification; reduced cortical bone ; reduced bone mineralization	No thyroid 2000 fold elevation of TSH and undetectable TH Functional TSHR	mansouri et al. (39); Friedrichsen et al. (40)
TSHR	Tshr deleted	Die unless treated with TH after weaning	Thyroid hypoplasia Undetectable T4 and T3, elevated TSH	Marians et al. (46)
Hyt/hyt	Loss of function mutation in the <i>Tshr</i> gene	Impaired linear growth, delayed endochondral ossification; reduced cortical bone ; reduced bone mineralization	2000 fold elevation of TSH and low TH Non-functional TSHR	Beamer et al. (42) ;Gu et al. (43)
D2	Deiodinase 2	Normal development and growth	Increase in T4 and TSH, normal T3	Bassett et al. (48)
TR α	α 1 and α 2 deleted Increase Δ α 1 but normal Δ α 2 and TR β expression	mice die within 5 weeks Severe growth retardation; delayed endochondral ossification; impaired chondrocyte differentiation; reduced mineralization	Severe hypothyroidism (T3 and T4 are 10% of WT levels); T3 injections rescue phenotype ; GH normal	Fraichard et al. (17)
TR α 1	α 1, Δ α 1 deleted Normal α 2, β 1 and β 2 expression	No growth retardation Not determined	Mild hypothyroidism in males but not females	Wikstrom et al. (122)
TR α 2	α 2, Δ α 2 deleted Increase in α 1 expression	No difference in the length of femur and tibia Normal endochondral ossification	Mild hypothyroidism Normal GH and low IGF-1	Salto et al. (123)
TR α ^{0/0}	α 1, Δ α 1, α 2, Δ α 2 deleted	Transient growth retardation; delayed endochondral ossification; impaired chondrocyte differentiation; reduced mineralization	Slight decrease in T4, normal TSH and T3; Normal GH and IGF-1	Gauthier et al. (52)
TR α 1 ^{PV/+}	Heterozygous dominant negative TR α receptor	Severe persistent growth retardation; delayed intramembraneous and endochondral ossification; impaired chondrocyte differentiation; reduced mineralization	Mild and transient systemic hypothyroidism Normal GH, reduced IGF-1	Kaneshige et al. (53)
TR β	TR β 1, β 2 deleted	Normal growth Persistent short stature; advanced endochondral and intramembraneous ossification; increased mineralization	RTH and goiter ;Increase in T4,T3 and TSH	Forrest et al. (5)
TR β 2	TR β 2 deleted Normal TR β 1, TR α expression	Normal growth Not determined	Mild RTH ;Increase in T4,T3 and TSH; 30% decrease in GH	Abel et al. (23)
TR β ^{PV/PV}	Homozygous dominant negative TR β receptor	Accelerated prenatal growth; persistent postnatal growth retardation; advanced intramembraneous and endochondral ossification; increased mineralization	Severe RTH and goiter ;400 fold elevation of TSH, and 15 fold elevation of T4 ; Reduced GH and increased IGF-1	Kaneshi et al. (59); O'shea et al. (57-58) ; Bassett et al. (56)
TR α ;TR β	α 1, α 2, β 1 and β 2 deleted	mice die within 5 weeks Disorganization of growth plate; delayed endochondral ossification; impaired chondrocyte differentiation; reduced mineralization	RTH and small goiter ;Increase in T3,T4 and TSH;	Gauthier et al. (111)

Cons.

KO	Deleted Protein	Skeletal Phenotype	TH-pituitary axis	Reference
TR α 1;TR β	α 1, $\Delta\alpha$ 1, β 1 and β 2 deleted	Persistent growth retardation; delayed endochondral ossification; reduced mineralization	RTH and large goiter ;Several fold increase in total and free T3, T4; increase in TSH ; reduced GH/IGF-1	Gothe et al. (124)
TR α 2;TR β	α 2 , $\Delta\alpha$ 2, β 1 and β 2 deleted	Transient growth delay	Mild hypothyroidism	Ng et al. (125)
TR $\alpha^{0/0}$;TR β	α 1, $\Delta\alpha$ 1, α 2 , $\Delta\alpha$ 2, β 1 and β 2 deleted	Disorganization of growth plate; growth delay; delayed endochondral ossification; impaired chondrocyte differentiation; reduced mineralization	RTH and goiter ;More than 10-fold increase in T4,T3 and TSH ;marked reduction in GH/IGF-1	Gauthier et al. (52)

TSH receptor (TSHRP556L) (41-43). Both mutants have a 2000-fold elevation of TSH and undetectable THs, but the TSHR is functional in Pax8^{-/-} mice whereas it is non-functional in hyt/hyt mice. Thus, the reciprocal relationship between THs and TSH remains intact in Pax8^{-/-} mice but is disrupted in hyt/hyt mice (44). Both mutants exhibited a similar skeletal phenotype of impaired linear growth, delayed endochondral ossification, impaired chondrocyte differentiation, reduced cortical bone, impaired trabecular bone remodeling and reduced bone mineralization (44-45). These data indicate that any action of TSH in bone is likely to be minor when compared to the effects of T3.

TSHR knockout mice (TSHR^{-/-}) present with developmental and growth delays and severe osteoporosis at all sites and profound hypothyroidism, with no detectable T4 and T3 and elevated TSH. The TSHR knockout mice die within 1 week of weaning unless treated with TH supplementation (46). TH replacement after weaning did not normalize the low bone mineral density in TSHR^{-/-} mice and these mice showed a dramatic increase in bone turnover. These findings confirmed that the bone loss was independent of T3 and T4 levels. Thus, an independent role for TSH in negative regulation of bone remodeling was proposed (47).

Deiodinase D2 knockout mice were generated to investigate the effects of osteoblast-specific T3 deficiency (48) because D2 expression was detected only in mature osteoblasts (12). The D2 knockout mice exhibit pituitary resistance to feedback regulation by T4 characterized by an increase in TSH and T4, but normal T3. Overall growth as well as skeletal development was found to be normal in D2 knockout mice indicating an insignificant role for D2 during endochondral and intramembranous ossification *in vivo*. However, adult D2 knockout mice had brittle bones due to increased mineralization and reduced bone formation because of restricted cellular hypothyroidism in osteoblasts (48). Further studies are needed to determine the mechanism

for reduced bone formation in the D2 conditional knockout mice.

Dual oxidase generates the hydrogen peroxide required by thyroid peroxidase for the incorporation of iodine into thyroglobulin. Dual oxidase 2 (Duox2) mutant mice, genetic model for congenital hypothyroidism, exhibit low T4 and high TSH caused by lack of T4 feedback to the pituitary gland and is associated with dwarfing and hearing impairment. Concomitant with small size and low aBMD, circulating levels of IGF-1 are lower in the mutants. These phenotypes of Duox2 mutant mice are consistent with lack of TH support, a key role for TH in skeletal development (49). Further studies are needed to investigate the specific functions of Duox2 in other tissues.

TR α mutants

Several TR α knockout mice have been generated and this has led to the identification of additional TR α isoforms expressed from a promoter within intron 7 of the THRA gene (50). As a result only TR $\alpha^{0/0}$ mice lack all TR α isoforms whereas other TR α mutants retain truncated isoforms with dominant-negative activity. TR $\alpha^{0/0}$ mice lack all TR α isoforms and are systemically euthyroid. During development TR $\alpha^{0/0}$ mice display features of skeletal hypothyroidism with transient growth retardation and with delayed endochondral ossification characterized by impaired chondrocyte differentiation and reduced mineralization (51-52).

Mice harboring dominant-negative mutations of TR α 1 in different genetic backgrounds have mild and transient systemic hypothyroidism but they also exhibit a more severe phenotype of delayed skeletal development than TR $\alpha^{0/0}$ mice. TR α 1^{PV/+} mice are most severely affected, displaying persistent post-natal growth retardation and markedly delayed endochondral ossification and decreased mineralization (38, 53).

Deletion or mutation of TR α does not affect systemic thyroid status but caused local skeletal hypothyroidism

while the presence of a dominant-negative TR α leads to a more severe skeletal phenotype than receptor deficiency alone. These findings demonstrate an important role for unliganded TR α 1 acting as a potent dominant-negative antagonist (54). The repressive actions of the unliganded receptor, therefore, have a greater physiologic effect than having no receptor at all. Consistent with this phenotype, skeletal expression of the T3 target genes, fibroblast growth factor receptors (FGFR) 1 and 3, were reduced (51, 55-56).

TR β mutants

Deletion of TR β results in elevated T4, T3 and TSH levels consistent with resistance to TH (RTH). In contrast to TR α mutants, during development TR β ^{-/-} mice display features of skeletal hyperthyroidism with advanced endochondral and intramembranous ossification, accelerated chondrocyte differentiation, increased mineralization and persistent short stature due to premature growth plate quiescence (5, 23, 51).

Mice harboring a dominant-negative mutation in THRB (TR β ^{PV/PV}) have severe RTH and exhibit a more severe phenotype than TR β ^{-/-} with accelerated prenatal growth characterized by advanced endochondral and intramembranous ossification (57-59). Consistent with phenotype, increased skeletal expression of the T3 target genes, FGFR1 and 3, (51, 55-56) revealed the presence of enhanced T3 action resulting from supraphysiological stimulation of TR α in bone. Thus, deletion or mutation of TR β disrupts the HPT axis resulting in skeletal thyrotoxicosis. The presence of a dominant negative TR β leads to a more severe skeletal phenotype than receptor deficiency alone. In summary, during development reduced T3 action in TR α mutant mice results in delayed ossification and reduced mineralization whereas increased T3 action in TR β mutant mice leads to advanced ossification and increased mineralization.

Clinical manifestation of TH signaling related mutations in humans (Table 2)

The developing skeleton is sensitive to thyroid status and childhood hypothyroidism is characterized by growth retardation, delayed bone age and short stature, whereas juvenile thyrotoxicosis accelerates growth and advances bone age but results in persistent short stature due to premature fusion of the epiphyses (2, 60-61). A loss of function mutation of the TSH β -subunit results in TSH deficiency and congenital hypothyroidism. Two affected siblings received TH replacement from birth but despite the lifelong absence of TSH, their skeletal development and bone mineral density were normal (62). These findings suggest that TSH is not required for normal skeletal

development and growth. TSHR mutations result in wide spectrum of clinical manifestations ranging from mild to severe hypothyroidism and hyperthyroidism (63). Up to this date more than 40 kinds of loss of function mutations in the TSHR gene have been reported as the causative defect in congenital hypothyroidism (64). By contrast, gain of function mutations in the TSHR gene were identified in familial non-autoimmune hyperthyroidism or sporadic non-autoimmune hyperthyroidism (65).

The syndrome of resistance to TH (RTH) is caused by decreased tissue responsiveness to TH and was first described in 1967 (66-67). Clinical features include goiter, elevated circulating TH levels, nonsuppressed serum TSH level, clinical euthyroidism, and tachycardia; some individuals also demonstrate attention deficit disorder and deficits of linear growth, hearing, and bone formation (66). The RTH genetic defect is caused mostly by mutations of TH receptor β gene. A dominant-negative mutation of TR β blunts negative feedback in the HPT axis and is characterized by elevated T4 and T3 and an inappropriately normal or elevated TSH. RTH patients display variable skeletal phenotypes that are difficult to interpret due to confounding effects of treatment and the heterogeneity of TR β mutations (66, 68). Nevertheless, reported features of increased bone turnover, osteoporosis, fracture and craniosynostosis in RTH suggest a predominant role for T3 in bone development (68). Last year, two families with a heterozygous mutation of THRA, resulting in expression of a dominant negative TR α protein, have been the first to be reported (69-70). Levels of free T4 and rT3 in these patients were in the low-normal range and T3 in the high-normal range, with normal TSH. Despite near normal TH levels, patients displayed a phenotype consistent with the characteristic features of hypothyroidism that included short stature, delayed bone development, skeletal dysplasia and chronic constipation, thus suggesting that TR α 1 plays a major role in human skeletal development. Recently, several patients with homozygous TR β mutations were reported (71). The clinical manifestations are a combination of those found in individuals heterozygous for a mutation only in TR α or in TR β . Patients have a more severe RTH phenotype, goiter, hearing loss, and much greater elevations of serum T3, T3, and TSH, than heterozygous individuals and also have neurological impairment and growth retardation, more characteristic of a deficiency in the action of TR α (69-70). These clinical phenotypes suggest the possible interference of the mutant TR β with function of TR α 1.

A sex-linked form of mental retardation with motor abnormalities, named the Allan-Herdon Dudley syndrome, was described in 1944. When MCT8, a specific trans-

Table 2 Clinical manifestation of TH signaling related mutations in humans

Clinical status	Mechanism	Clinical manifestation	TH-pituitary axis	Reference
Congenital hypothyroidism	Loss of function TSHR mutation	Prolonged jaundice, constipation, poor feeding, umbilical hernia, macroglossia, wide open posterior fontanel and edematous and dry skin	Mild to severe hypothyroidism or thyroid hypoplasia	Papadimitriou et al. (62); Persani et al. (64)
Resistance to Thyroid Hormone	Heterozygous dominant negative TR α receptor	Short stature; delayed bone development; transient delay in motor development; mild impairment of cognitive development; chronic constipation	Low normal free T4 and rT3; High T3 ; Normal TSH	Boucukova et al. (69);van Mullem et al. (70)
Resistance to Thyroid Hormone	Homozygous dominant negative TR β receptor	Large goiter; dysmorphic features; severe tachycardia; developmental and growth delay; mental retardation; hearing deficit	High T4 and T3; nonsuppressed TSH	Ferrara et al. (71)
	Heterozygous dominant negative TR β receptor	Variable, can induce goiter; hearing deficit; hyperactive behavior; learning disability; developmental delay; tachycardia	High T4 and T3; nonsuppressed TSH	Refetoff et al. (66-67); Weiss et al. (126)
TH cell transporter defect (THCTD)	MCT8 mutation	Hypotonia, motor delay, feeding problem, no speech development, spasticity, cognitive impairment, normal linear growth	High serum T3 and low rT3 and T4; Normal or slightly elevated TSH	Friesema et al. (72); Dumitrescu et al. (73)
TH metabolism defect (THMD)	SECISBP2 mutation	Short stature , delayed bone age in child Transient growth retardation	High T4 and rT3; Low T3; Slightly elevated TSH.	Dumitrescu et al. (74); Refetoff et al. (66)

porter of TH, was sequenced in these patients, inactivating mutations were identified in some individuals (72-73). Patients manifest with truncal hypotonia, poor head control, and later spasticity and were found to have abnormal thyroid function (elevated serum T4 and rT3 and low T3).

Deiodinases are selenoproteins that catalyze iodothyronine deiodination and are important in TH activation and inactivation. Selenium is an essential trace element required for the biosynthesis of selenoproteins and selenocysteine insertion sequence binding protein 2 (SECISBP2), a key trans-acting factor. Patients with mutations in the *SECISBP2* gene presented with transient growth retardation associated with abnormal thyroid function, low T3, high T4 and rT3, and slightly elevated TSH (74). Thus, there are a number of clinical genetic studies with mutations in genes related to the TH signaling pathway that attest the importance of TH in the regulation of skeletal metabolism.

Effects of T3 in bone cells in vitro

Chondrocytes

TR α 1 and TR β 1 are expressed in resting and proliferating chondrocytes in the growth plate, suggesting these cells

are direct targets for T3 actions (75). T3 stimulates clonal expansion of resting chondrocyte progenitor cells but inhibits subsequent chondrocyte proliferation, while stimulating hypertrophic differentiation (75-79). Accordingly, T3 induces markers of hypertrophic chondrocyte differentiation, including alkaline phosphatase and collagen X expression in primary growth-plate chondrocyte cultures, and enhances cartilage matrix mineralization (75). T3 also stimulates the expression of proteoglycan and collagen-degrading enzymes including aggrecanase-2 (a disintegrin and metalloproteinase with thrombospondin motifs1, ADAMTS5) and matrix metalloproteinase 13 (MMP13) (78, 80-81). T3 regulation of growth plate chondrocyte proliferation and differentiation *in vitro* has been shown to involve a number of growth factor signaling pathways including IGF-1, Wnt, the Ihh/PTHrP feedback loop and FGFR3 (Figure 2) (55, 82-83). In conclusion, TH stimulates maturation of chondrocytes and the progression of endochondral ossification and is essential for linear growth.

Osteoblasts

T3 has been found to stimulate, inhibit, or exert no effect on osteoblastic cell proliferation, but a consensus

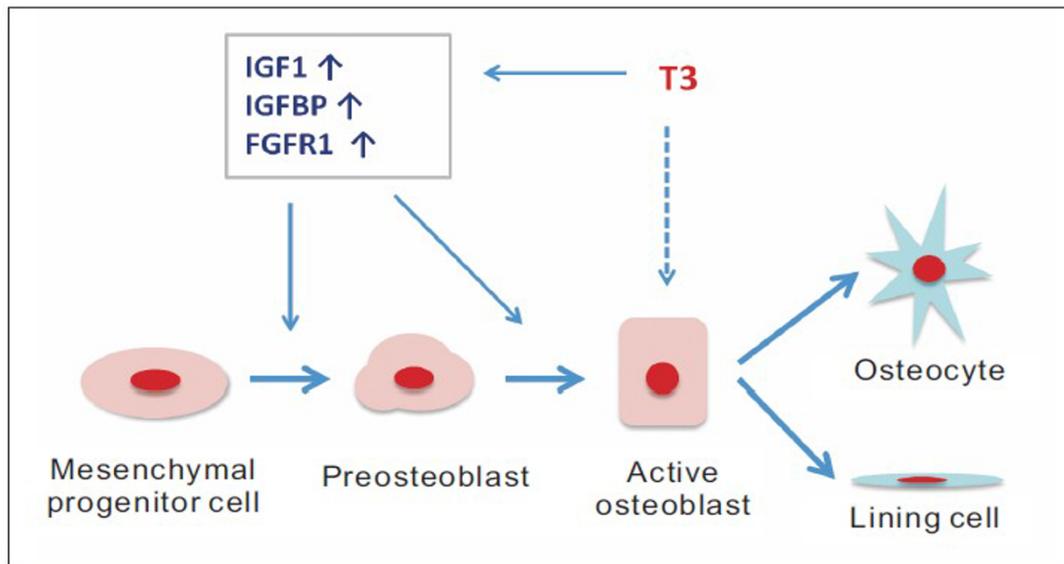


Figure 3 Mechanisms for T3 regulation of osteoblast differentiation. T3 increases local IGF-I actions by modulating production of IGF-I and/or its binding proteins and thereby stimulate differentiation of mesenchymal cells into osteoblast lineage. T3 can also stimulate local FGF actions by increasing FGF receptor expression in osteoblasts. T3 can also exert direct effects via TRE to regulate transcription of bone formation genes.

it mediates both GH-dependent and -independent effects and is involved in chondrocyte recruitment, proliferation and hypertrophic differentiation (103).

IGF-1 signaling is also required to maintain the *Ihh*-PTHrP loop during skeletogenesis. In fetal *IGF1^{-/-}* mice, expression of *Ihh* was reduced in long bones, whereas expression of PTHrP was increased (104). Furthermore, Wang *et al* (105) have demonstrated that IGF-1/IGF1R induces *Wnt4* expression and β -catenin activation. IGF-1/IGF1R actions on growth plate chondrocytes are neutralized by the Wnt antagonists sFRP3 and Dkk1, confirming that the *Wnt*/ β -catenin signaling pathway is downstream of IGF-1 signaling (Figure 2).

It is now known that IGF-1 expression in bone is regulated by GH as well as many systemic and local regulators of bone growth. In this regard, our previous studies showed that days 7 to 14 of the prepubertal period and days 23 to 31 of the pubertal period represent critical time points for skeletal growth as well as increases in serum IGF-I levels in mice (106). It was also determined that there is an important critical period during prepubertal growth when the effects of GH are small and IGF-I remains an important regulator (101). Recently, we demonstrated that TH is a major regulator of IGF-I expression, independent of GH, during the prepubertal growth period (45). When we measured the postnatal changes in serum levels of total T3 and IGF-1 in C57BL/6J mice, as seen in Figure 4, serum levels of IGF-1 increased almost three fold in mice during the prepubertal growth period. This increase in IGF-1 was preceded

by changes in serum T3 levels, which increased two-fold between day 7 and 14. Studies using genetic mouse models deficient in TH found that serum levels of IGF-1 were reduced by more than 50% at day 21 compared to wild-type mice as a consequence of a decrease in IGF-1 expression in liver and bone. Daily administration of T3/T4 during the prepubertal growth period (days 5 to 14) increased IGF-1 expression in both liver and bone and normalized the serum IGF-1 levels. Furthermore, *in vitro* studies in osteoblasts revealed that TH, in the presence of TR α 1, bound to a TH response element in intron 1 of the IGF-1 gene to stimulate transcription (45). Thus, TH is a key regulator of both local and endocrine IGF-1 action during the prepubertal growth period.

In terms of the target cell types for TH effects on IGF-1 expression, we and others have found that TH treatment increased IGF-1 expression in bone cells and chondrocytes (45, 58, 107). There is also evidence that TH is able to act directly on growth plate chondrocytes through GH-independent mechanisms (108). Lewinson *et al* demonstrated that TH is able to stimulate longitudinal bone growth in animals in which GH secretion has been ablated by hypophysectomy (109). TH also stimulated chondrocyte differentiation in both thyroidectomized and hypophysectomized rats, a role for which growth hormone cannot substitute (110). Thus, there is ample evidence that TH is essential for skeletal growth during the prepubertal growth period.

The relationship between TH signaling and GH/IGF-1 signaling is potentially complex due to multiple possible

points of interaction at both the systemic and local levels. Through studies using a variety of transgenic mice in which TR function has been altered, our understanding of this complexity has been improved. $TR\alpha^{0/0}$ and $TR\alpha^{PV/+}$ mice showed normal pituitary GH production, but diminished expression of GH receptor, IGF-1, and IGF-1 receptor in growth plate chondrocytes (52, 58). By contrast, pituitary GH production was reduced and GH receptor and IGF-1 expression in the growth plate was increased in $TR\beta^{-/-}$ and $TR\beta^{PV/PV}$ mice (5, 58, 111). The different phenotypes between the $TR\alpha^{PV/+}$ mice and $TR\beta^{PV/PV}$ mice can be explained by the known finding that the pituitary gland is a $TR\beta$ -responsive tissue, while the growth plate is primarily a $TR\alpha$ -responsive tissue. Thus, GH/IGF1 signaling is also a local downstream mediator of T3 action in the growth plate.

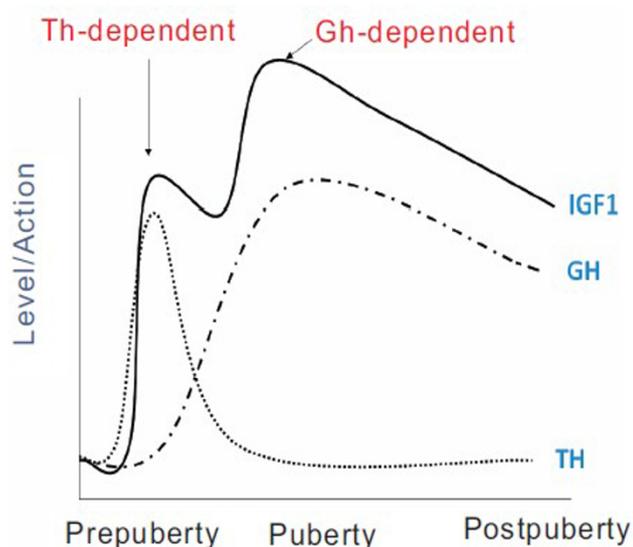


Figure 4 Model of TH regulation of skeletal growth in mice. It is proposed that the increase in IGF-I expression and bone accretion during prepubertal and pubertal growth periods are mediated via TH and GH-dependent mechanisms, respectively. Some of the effects of TH on bone growth are direct and occur independent of GH.

GH therapy normalized the decreased serum IGF-1 level of hypothyroid pups. However, growth failure, despite normal serum IGF-1 concentrations in GH-treated hypothyroid rats, has been described earlier (112). Consistent with this, we found that inhibition of IGF-1 action only partially reduced TH effects on osteoblast differentiation and the rate of metatarsal bone mineralization (45). These results provide indirect evidence that TH acts via other pathways, in addition to IGF-1. It is known that TH influences fibroblast growth factor (FGF) receptor signaling in bone and interacts with the Wnt/ β -catenin and *Ihh*/PTHrP signaling pathways to regulate endo-

chondral ossification (52–54). The extent to which IGF-1 interaction with other growth factor signaling pathways is involved in TH regulation of skeletal metabolism remains to be determined.

*Interaction with the *Ihh*/PTHrP feedback loop*

The pace of chondrocyte differentiation is precisely regulated by the *Ihh*/PTHrP paracrine negative feedback loop. Prehypertrophic chondrocytes secrete *Ihh* which diffuses to periarticular cells to induce synthesis of PTHrP. PTHrP, acting via its receptor PTHR1, then completes the loop by stimulating chondrocyte proliferation and inhibiting further hypertrophic differentiation (113–114). The effect of TH on the *Ihh*/PTHrP feedback loop has been studied in thyroid manipulated rats (83). In hypothyroid rats, growth plates were grossly disorganized and the expression of PTHrP was increased and extended throughout the growth plate, while the PTHrP receptor was expressed in the same location as in euthyroid animals. These changes in PTHrP receptor expression can lead to inhibition of hypertrophic chondrocyte differentiation and result in arrest of linear growth (Figure 2). In thyrotoxic growth plates, histology essentially was normal but PTHrP receptor (PTHrP-R) mRNA was undetectable, while PTHrP mRNA expression was unchanged. An absence of the PTHrP receptor results in negative PTHrP signaling and progression of hypertrophic chondrocyte differentiation to accelerate linear growth. Furthermore, recent studies in chicken tibia explants have shown that *Ihh* stimulates degradation of the type 2 deiodinase enzyme resulting in an induction of PTHrP expression (114). Together, these findings suggest that TH regulates the set point of the *Ihh*/PTHrP feedback loop to modulate the pace of chondrocyte differentiation and endochondral bone formation during postnatal growth.

Interaction with Fibroblast Growth Factor signaling

Signaling through the FGF pathway has been demonstrated to negatively regulate proliferation of growth plate chondrocytes. During endochondral ossification, FGFR2 is expressed in condensing mesenchyme and perichondrium, FGFR1 is present in prehypertrophic and hypertrophic chondrocytes, and proliferating chondrocytes express FGFR3. During linear growth, FGFR2 is not expressed in the growth plate, but FGFR1 expression persists in prehypertrophic and hypertrophic cells, and proliferating chondrocytes express FGFR3 (33, 115). The role of FGFR1 and FGFR2 in the developing growth plate is poorly understood. However, FGFR3 has been predicted to play an important role. Activating mutations in FGFR3 results in achondroplasia, the most common form of dwarfism in humans (115–116), whereas *Fgfr3* knockout

mice display limb overgrowth (117-118), indicating that FGFR3 is a negative regulator of linear growth. T3 stimulates expression of FGFR-1 and FGFR-2 mRNA in the ATDC5 chondrocytic cell line undergoing chondrogenesis, but stimulation of FGFR3 by T3 was greater and persisted longer, coinciding with the period in which T3 inhibited chondrocyte proliferation and stimulated hypertrophic differentiation (55). Investigation of the FGF/FGFR signaling pathway in TR mutant mice revealed that FGFR 3 was reduced in growth plates of TR $\alpha^{D/D}$ and TR $\alpha^{PV/+}$ mice and increased in TR $\beta^{-/-}$ and TR $\beta^{PV/PV}$ mice (55-58). Taken together, these findings suggest that FGFR3 may play a role in mediating effects of T3 on chondrogenesis. Some of the growth inhibitory actions of FGFR3 are mediated via reduced activity of the Ihh/PTHrP feedback loop (119). The interactions between FGFR3 and other growth factor signaling pathways in regulating chondrocytes still need to be worked out.

Interaction with Wnt/ β -catenin signaling

Wnt/ β -Catenin signaling also has been recognized as an important signal-transduction pathway in regulating terminal differentiation of growth plate chondrocytes. Inhibition of β -catenin signaling in Col2a1-ICAT transgenic mice results in reduced chondrocyte proliferation and differentiation, delayed formation of the secondary ossification center, and reduced skeletal growth (120). Wang *et al* have shown that T3 promotes growth plate chondrocyte terminal differentiation by increasing Wnt-4 expression, β -catenin accumulation, TCF/LEF transcriptional activity, and expression of the Wnt/ β -catenin target gene *Runx2/cbfa1* (82). Furthermore, they also have shown that T3 treatment stimulates PI3K/Akt/GSK-3 β signaling (105). PI3K and Akt are important signal transducers of IGF-1 signaling. Akt can inactivate GSK-3 β , a negative regulator of the canonical Wnt/ β -catenin pathway (121). The inhibition of PI3K/Akt activity by LY294002 prevents T3-induced Wnt4 expression and β -catenin activation. These results indicate that TH promotes growth plate cell differentiation and longitudinal bone growth by activating β -catenin signaling via modulation of IGF-1/IGF1R signaling through the Wnt and PI3K/Akt pathways. While a number of growth factor signaling pathways have been implicated to play an important role in the TH regulation of skeletal growth (Figure 3), the extent to which these growth factor signaling pathways contribute to TH action *in vivo* remains to be determined.

Conclusions

There is now considerable data in the literature both

from mouse genetic studies and human clinical studies involving mutations in genes related to the TH signaling pathway that demonstrate a key role for TH in the regulation of skeletal growth. In terms of target cell types for TH action, while much focus is on chondrocytes, there is also recent evidence that other bone cell types including osteoblasts and osteoclasts are also regulated by TH signaling. In terms of the mechanism for TH action, studies suggest that TH regulates a number of key growth factor signaling pathways including IGF-1, Wnt, PTHrP and FGF to regulate skeletal growth. However, the relative contribution of these growth factor signaling pathways in mediating TH effects on bone *in vivo* remain to be determined. The issue of how the various growth factor signaling pathways interact to mediate TH effects in various target cell types needs to be investigated. Also, studies are needed to address the issue of whether all of the TH effects on bone cells are mediated via genomic actions of TH or some of the TH effects are mediated via non-genomic actions. Future development and application of mice with conditional disruption of genes involved in TH signaling pathway in time and space are required to identify the cell-specific mechanisms of TH action and interaction with various pathways *in vivo*.

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