

## REVIEW

# Insight into the physiological actions of thyroid hormone receptors from genetically modified mice

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### Introduction

Thyroid hormones exert a range of developmental and physiological actions in all vertebrates. Serum concentrations of L-thyroxine (T4) and 3,5,3'-L-tri-iodothyronine (T3) are maintained by a negative feedback loop involving T3 inhibition of hypothalamic thyrotrophin-releasing hormone (TRH) and pituitary thyroid-stimulating hormone (TSH) secretion, and by tissue-specific and hormone-regulated expression of the three iodothyronine deiodinase enzymes that activate or metabolise thyroid hormones (Bianco *et al.* 2002). T3 actions are mediated by two T3 receptors, TR $\alpha$  and TR $\beta$ , which act as hormone-inducible transcription factors. The TR $\alpha$  (NR1A1) and TR $\beta$  (NR1A2) genes encode mRNAs that are alternatively spliced to generate nine mRNA isoforms (TR $\alpha$ 1,  $\alpha$ 2,  $\alpha$ 3,  $\Delta\alpha$ 1,  $\Delta\alpha$ 2,  $\beta$ 1,  $\beta$ 2,  $\beta$ 3 and  $\Delta\beta$ 3), of which four (TR $\alpha$ 1,  $\alpha$ 2,  $\beta$ 1 and  $\beta$ 2) are known to be expressed at the protein level *in vivo* (Lazar 1993, Chassande *et al.* 1997, Williams 2000) (Fig. 1). The numerous TR mRNAs are expressed widely in tissue- and developmental stage-specific patterns, although it is important to note that levels of mRNA expression may not correlate with receptor protein concentrations in individual tissues (Forrest *et al.* 1990, Schwartz *et al.* 1994). The TR $\alpha$ 2,  $\alpha$ 3,  $\Delta\alpha$ 1 and  $\Delta\alpha$ 2 transcripts encode proteins that fail to bind T3 *in vitro*. These 'non-binding' isoforms, in addition to TR $\Delta\beta$ 3 which does bind hormone, may act as dominant negative antagonists of the true T3-binding receptors *in vitro*, but their physiological functions and those of the TR $\beta$ 3 isoform have not been determined. In order to obtain a new understanding of the complexities of T3 action *in vivo* and the role of TRs during development, many mouse models of disrupted or augmented thyroid hormone signalling have been generated. The aim of this review is to provide a picture of the physiological actions of thyroid hormones by considering the phenotypes of these genetically modified mice.

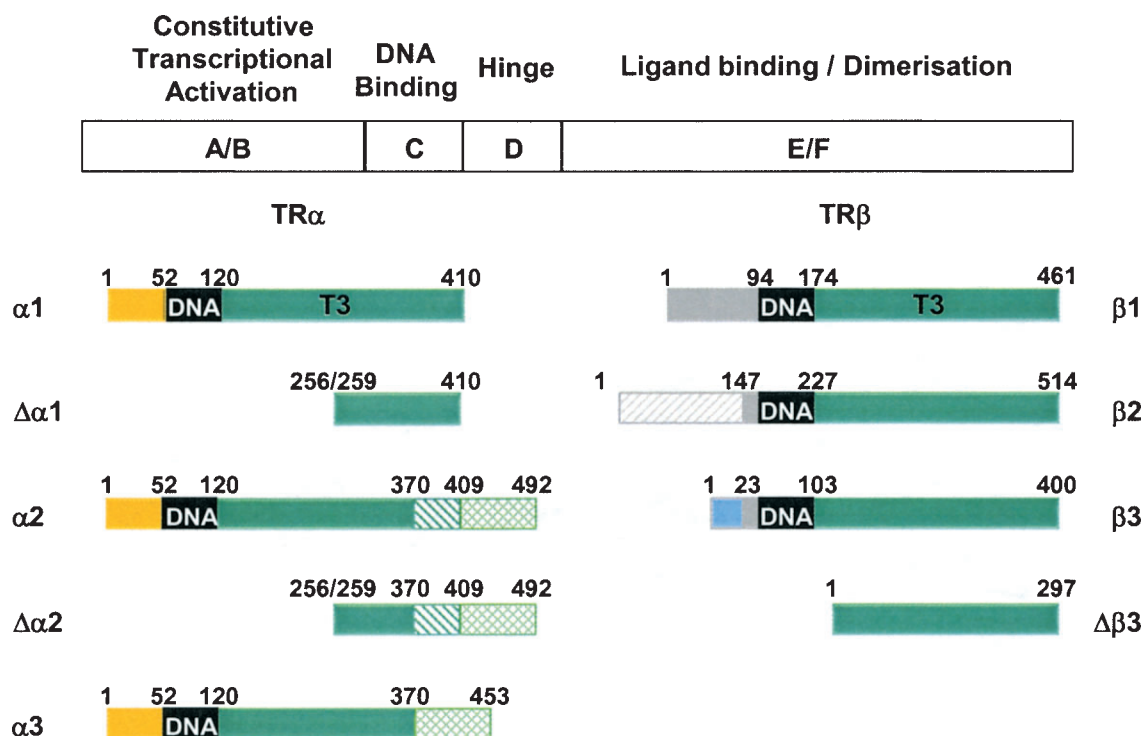
### Endocrine system

#### TR $\alpha^{0/0}$ knockout

TR $\alpha^{0/0}$  mice lack expression of all products of the TR $\alpha$  locus (Gauthier *et al.* 2001) (Table 1). Animals survive normally and are fertile with normal basal thyroid status and pituitary growth hormone (GH) mRNA concentrations (Gauthier *et al.* 2001). Nevertheless, compared with wild-type littermates, TR $\alpha^{0/0}$  mice display increased sensitivity to thyroid hormones in the pituitary and liver following provocative testing with increasing doses of T3 to suppress TSH and T4 secretion (Macchia *et al.* 2001).

#### TR $\alpha^{-/-}$ knockout

Disruption of exon 2, the first coding exon within the TR $\alpha$  locus, prevents transcription of both TR $\alpha$ 1 and TR $\alpha$ 2 mRNAs, but does not interfere with expression of TR $\Delta\alpha$ 1 and  $\Delta\alpha$ 2 from an internal promoter in intron 7 (Chassande *et al.* 1997, Fraichard *et al.* 1997) (Table 1). TR $\alpha^{-/-}$  pups die shortly after weaning unless treated with T3 (Fraichard *et al.* 1997). The thyroid gland develops abnormally and homozygous mutants are profoundly hypothyroid with T3 and T4 concentrations reduced to 40% and <10% of wild-type concentrations by 5 weeks of age. The greater reduction in T4 compared with T3 suggests that progressive hypothyroidism results from defective hormone production rather than an abnormality in peripheral T3 production (Fraichard *et al.* 1997). Accordingly, pituitary TSH $\beta$  mRNA expression was decreased more than 3-fold in TR $\alpha^{-/-}$  animals although GH mRNA levels were unaffected. Homozygous mutants die within 5 weeks, although they can be rescued by hormone replacement for only 1 week (1  $\mu$ g T3 subcutaneous injection) during the third week of life at the time of weaning. Thereafter, animals gain weight and hormone levels of 2.5-month-old rescued animals are normal



**Figure 1** Thyroid hormone receptor isoforms. Protein products arising from the TR $\alpha$  and TR $\beta$  genes are displayed below a schematic diagram of the functional domains present within proteins of the nuclear receptor superfamily. Numbering represents the amino acid positions within the rat TR proteins. Shading indicates identical shared regions and differences between TR variants that are translated from alternately spliced mRNA isoforms.

**Table 1** Genotypes and thyroid status of TR null and related knockout mice

	Reference	Deleted TR mRNAs	Expressed TR mRNAs	Thyroid status
<b>Knockout</b>				
TR $\alpha$ <sup>-/-</sup>	Fraichard <i>et al.</i> (1997)	$\alpha 1$ , $\alpha 2$	$\Delta\alpha 1$ , $\Delta\alpha 2$ , all $\beta$ isoforms	Progressive to severely hypothyroid
TR $\alpha 1$ <sup>-/-</sup>	Wikstrom <i>et al.</i> (1998)	$\alpha 1$ , $\Delta\alpha 1$	$\alpha 2$ , $\Delta\alpha 2$ , all $\beta$ isoforms	Mildly hypothyroid
TR $\alpha 2$ <sup>-/-</sup>	Salto <i>et al.</i> (2001)	$\alpha 2$ , $\Delta\alpha 2$	$\alpha 1$ (over-expressed), $\Delta\alpha 1$ , all $\beta$ isoforms	Mixed hyper/hypothyroid
TR $\alpha$ <sup>0/0</sup>	Gauthier <i>et al.</i> (2001)	All $\alpha$ isoforms	All $\beta$ isoforms	Euthyroid
TR $\beta$ <sup>-/-</sup>	Forrest <i>et al.</i> (1996b)	All $\beta$ isoforms	All $\alpha$ isoforms	RTH
TR $\beta 2$ <sup>-/-</sup>	Abel <i>et al.</i> (1999a), Ng <i>et al.</i> (2001)	$\beta 2$	$\beta 1$ , $\beta 3$ , $\Delta\beta 3$ , all $\alpha$ isoforms	RTH
TR $\alpha$ <sup>0/0</sup> $\beta$ <sup>-/-</sup>	Gauthier <i>et al.</i> (2001)	All $\alpha$ isoforms, all $\beta$ isoforms	—	Profound RTH
Pax8 <sup>-/-</sup>	Mansouri <i>et al.</i> (1998)	Deficiency of thyroid follicular cells following thyroid gland agenesis		Severely hypothyroid

RTH indicates mice with resistance to thyroid hormone.

(Fraichard *et al.* 1997), although both male and female TR $\alpha$ <sup>-/-</sup> mice are infertile. These data indicate a stimulatory role for TR $\alpha 1$  or  $\alpha 2$  in the initiation of thyroid

hormone production at weaning or a suppressive role for the truncated  $\Delta\alpha 1$  and  $\Delta\alpha 2$  products that continue to be expressed in TR $\alpha$ <sup>-/-</sup> mice.

*TRα1<sup>-/-</sup> knockout*

Both male and female *TRα1<sup>-/-</sup>* mice are fertile and display mild abnormalities of the pituitary–thyroid axis. Male *TRα1<sup>-/-</sup>* mice have reduced free T4 concentrations but normal free T3 concentrations compared with wild-type mice whereas females are euthyroid. Thyroid gland histology in *TRα<sup>-/-</sup>* mice was normal. Reduced T4 levels in males occurred in association with a small reduction in TSH concentrations and reduced pituitary *TSHα*, but not *TSHβ*, mRNA expression. These data suggest that the mild hypothyroidism in male *TRα1<sup>-/-</sup>* mice results from impaired *TSHα* gene regulation (Wikstrom *et al.* 1998). In combination with findings in *TRβ<sup>-/-</sup>* mice (Forrest *et al.* 1996a,b), these studies suggest a model in which TSH gene expression is dynamically regulated by interactions between individual isoforms expressed by both *TRα* and *β* genes (Wikstrom *et al.* 1998).

*TRα2<sup>-/-</sup> knockout*

*TRα2<sup>-/-</sup>* mice overexpress *TRα1* as an inevitable consequence of gene targeting and are hypothyroid (Salto *et al.* 2001). Free T3 and T4 concentrations are reduced, but TSH concentrations are inappropriately normal and there is impaired hormone production from the thyroid gland. Nevertheless, there are also features of hyperthyroidism such as increased heart rate, weight loss and elevated body temperature, suggesting that tissue-specific differences in thyroid hormone responsiveness may depend on the amount of *TRα1* expressed. Thus, the *TRα1:α2* ratio may regulate hormone responsiveness in specific tissues. Female *TRα2<sup>-/-</sup>* mice have a prolonged oestrous cycle due to abnormal patterns of ovulation that are reflected by impaired fertility. In contrast, the *TRα2<sup>-/-</sup>* mice have normal GH levels but there is a decrease in circulating insulin-like growth factor-I (IGF-I) concentrations, suggesting that GH signalling requires tissue euthyroidism.

*TRβ<sup>-/-</sup> knockout*

*TRβ<sup>-/-</sup>* mice, which lack expression of all *TRβ* isoforms (Table 1), display resistance to thyroid hormone (RTH) and demonstrate the key role for *TRβ* in set-point control of the pituitary–thyroid feedback axis (Forrest *et al.* 1996b, Gauthier *et al.* 1999). There is enlargement of the thyroid gland with an increased size and number of follicles. The inappropriate elevation of TSH is accompanied by selective 2.5- and 3.3-fold rises in pituitary *TSHα* and *TSHβ* mRNA expression, suggesting that RTH results from defective thyrotroph function rather than hyperplasia or pituitary gland malformation (Forrest *et al.* 1996b). Following the suggestion that *TRβ* may regulate TSH gene transcription directly (Forrest *et al.*

1996b), further studies (Weiss *et al.* 1997) revealed that *TRβ* is not required for increased TSH expression in hypothyroidism and is not essential for T3-mediated repression of TSH. Thus, *TRβ* enhances the sensitivity of the system and is necessary for complete inhibition of TSH, thereby resulting in the RTH phenotype in *TRβ<sup>-/-</sup>* mice (Weiss *et al.* 1997). These data are in accord with previous studies *in vitro*, which indicate that *TRH* is negatively regulated by T3 acting via *TRβ* (Lezoualc'h *et al.* 1992) and suggest that *TRβ*-mediated control of the pituitary–thyroid axis operates at the level of the hypothalamus as well as the pituitary. Although *TRβ<sup>-/-</sup>* mice display features of RTH similar to those observed in humans, they provide a recessive model for this condition whereas RTH in man is autosomal dominant (Forrest *et al.* 1996a,b). *TRβ<sup>-/-</sup>* mice also exhibit impaired T3-dependent regulation of cholesterol metabolism (Gullberg *et al.* 2000, Macchia *et al.* 2001), supporting previous observations showing that basal cholesterol levels were elevated in *TRβ<sup>-/-</sup>* mice, despite elevated basal T4 concentrations (Weiss *et al.* 1998), thus demonstrating peripheral as well as pituitary RTH.

*TRβ2<sup>-/-</sup> knockout*

To elucidate further the role of *TRβ* in pituitary–thyroid feedback control, *TRβ2<sup>-/-</sup>* mice were generated (Abel *et al.* 1999a) (Table 1). These mice also have RTH with a 3-fold increase in basal T4 concentrations, a 40% increase in T3, a 2.5-fold elevation in TSH and defective TSH suppression following T3 administration. Pituitary *TSHβ* mRNA was increased in *TRβ2<sup>-/-</sup>* mice and there was impaired stimulation of pituitary GH mRNA following T3 administration, adding to previous evidence indicating that *TRβ2* plays a major role in T3-stimulated GH mRNA expression (Ball *et al.* 1997). Thus, deletion of *TRβ2* is sufficient to cause RTH, suggesting that *TRβ1* cannot adequately substitute for *β2* and that *β2* primarily mediates feedback control. This hypothesis is supported by studies which indicate that *TRβ2* is a more potent regulator of TSH mRNA than *β1* (Ng *et al.* 1995, Safer *et al.* 1997). More recently, Ng *et al.* (2001) developed *Thrb<sup>tm2Df/tm2Df</sup>* mutant mice, which also lack expression of *TRβ2*. These mice display milder RTH than observed by the same group previously in *TRβ<sup>-/-</sup>* mice (Forrest *et al.* 1996b) and milder than observed in *TRβ2<sup>-/-</sup>* mice reported by Abel *et al.* (1999a). Thus, Ng *et al.* (2001) concluded that *β2* acts co-operatively with *β1* to regulate the pituitary–thyroid axis, in contrast to conclusions from the studies discussed above that suggest that *β2* is the primarily important isoform (Abel *et al.* 1999a). Although the two laboratories each report mice that lack *TRβ2*, their phenotypic differences may result from the differing gene targeting strategies employed by the two groups or from differences in genetic background of the mice. Thus, it is difficult to conclude at this stage precisely how much

contribution to control of the pituitary–thyroid axis is made by TR $\beta$ 2 relative to  $\beta$ 1 *in vivo* or whether other modifier genes are influential.

#### TR $\alpha^{0/0}\beta^{-/-}$ double knockout

Deletion of all TR isoforms in TR $\alpha^{0/0}\beta^{-/-}$  mice (Gauthier *et al.* 2001) results in profound abnormalities of the pituitary–thyroid axis with concentrations of T4, T3 and TSH elevated 14-fold, 13-fold and more than 200-fold respectively, which are similar to findings in TR $\alpha 1^{-/-}\beta^{-/-}$  (Gothe *et al.* 1999) and TR $\alpha^{-/-}\beta^{-/-}$  mice (Gauthier *et al.* 1999). Pituitary GH mRNA expression is markedly reduced in TR $\alpha^{0/0}\beta^{-/-}$  double mutant mice in contrast to TR $\alpha^{0/0}$  animals, in which expression is preserved (Gauthier *et al.* 2001), providing further evidence that T3 regulation of GH occurs via TR $\beta$ .

The high levels of TSH in TR $\alpha^{0/0}\beta^{-/-}$  mice are around half those reported in Pax8 $^{-/-}$  mice (Flamant *et al.* 2002) in which the primary defect is an absence of thyroid follicular cells (Mansouri *et al.* 1998) (Table 1). The number of pituitary thyrotrophs is markedly elevated in both Pax8 $^{-/-}$  and TR $\alpha^{0/0}\beta^{-/-}$  mice and TSH $\alpha$  gene expression is similar in both genotypes (Flamant *et al.* 2002). These similar features occur despite the fact that Pax8 $^{-/-}$  mice have no circulating T3 and T4 whilst TR $\alpha^{0/0}\beta^{-/-}$  mice have markedly elevated peripheral hormone concentrations and suggest that programming of the numbers of developing pituitary thyrotrophs occurs independent of thyroid hormones or TRs.

We have selected Pax8 $^{-/-}$  mice (Mansouri *et al.* 1998) as the best available genetic model of isolated thyroid hormone deficiency for comparison with mice harbouring TR deletions. Whilst it remains possible that Pax8 has additional and uncharacterised functions in tissues other than the thyroid gland, the phenotype of Pax8 $^{-/-}$  mice appears to result solely from hypothyroidism. Furthermore, Pax8 $^{-/-}$  mice have been crossed with various TR-deficient mice (Flamant *et al.* 2002) enabling objective phenotypic comparisons to be made with TR-null mice. Nevertheless, alternative genetic models of hypothyroidism including the Ames (*df/df*) and Snell (*dw/dw*) dwarf and *hyt/hyt* (Beamer *et al.* 1981) hypothyroid mutants have been reported. Whilst these mice provide further useful tools for the further understanding of T3 action, the *df/df* and *dw/dw* mutants develop hypothyroidism secondary to hypopituitarism resulting from mutations of the prophet of Pit-1 (Sornson *et al.* 1996) and Pit-1 (Li *et al.* 1990) transcription factors respectively, whilst congenital hypothyroidism in *hyt/hyt* mice results from a mutation of the TSH receptor (Stein *et al.* 1994), which is expressed widely. These complexities suggest that comparison of the phenotypes of TR-null mice with the *df/df*, *dw/dw* and *hyt/hyt* mutants should be viewed cautiously. For the purposes of this review,

therefore, we have focused on Pax8 $^{-/-}$  mice as a standard genetic model of hypothyroidism.

Pax8 $^{-/-}$  mice have a complete absence of T4 in postnatal life, when autonomous production of thyroid hormone replaces the maternal supply, and survival is not seen beyond postnatal day 30 (Flamant *et al.* 2002). However, lethality following weaning is not necessarily a direct consequence of hypothyroidism. Studies of Pax8 $^{-/-}$ TR $\alpha^{0/0}$  and Pax8 $^{-/-}$ TR $\beta^{-/-}$  mutants revealed that Pax8 $^{-/-}$ TR $\alpha^{0/0}$  were viable and similar in phenotype to TR $\alpha^{0/0}\beta^{-/-}$  double receptor mutants, whilst the Pax8 $^{-/-}$ TR $\beta^{-/-}$  mutation was lethal and similar to Pax8 $^{-/-}$  single mutants. These data indicate detrimental developmental effects of unliganded TR $\alpha$  receptor isoforms in Pax8 $^{-/-}$  mice (Flamant *et al.* 2002) and support similar conclusions derived from TR $\alpha^{-/-}$  mice (Fraichard *et al.* 1997).

#### TR $\alpha^{-/-}\beta^{-/-}$ double knockout

In TR $\alpha^{-/-}\beta^{-/-}$  mice, T4 and T3 levels were elevated 10-fold and TSH increased more than 100-fold (Gauthier *et al.* 1999). There was concomitant thyroid gland enlargement with follicular cell hyperplasia, an increase in the number of follicles, which were smaller than seen in TR $\beta^{-/-}$  mice, and evidence of abnormal colloid production and increased vascularity of the gland (Gauthier *et al.* 1999). As in the case of TR $\alpha^{-/-}$  animals (Fraichard *et al.* 1997), TR $\alpha^{-/-}\beta^{-/-}$  mice die following weaning (Gauthier *et al.* 1999), further implicating the TR $\Delta\alpha 1$  and  $\Delta\alpha 2$  isoforms in the generation of this phenotype.

#### TR $\alpha 1^{-/-}\beta^{-/-}$ double knockout

Gothe *et al.* (1999) reported the phenotype of TR $\alpha 1^{-/-}\beta^{-/-}$  mice. Free T3 and T4 concentrations were elevated 60-fold with a 160-fold increase in TSH in 4- to 8-week-old mice that reduced to a 60-fold elevation at 5 months of age. There was a 15-fold increase in thyroid gland weight with hyperplasia and proliferation of follicular cells leading to progressive goitre. TSH $\alpha$  and TSH $\beta$  mRNA levels were increased 3.3- and 26-fold respectively in double mutant pituitary glands, although the pituitary gland was not malformed or enlarged. Similar to Pax8 $^{-/-}$  and TR $\alpha^{0/0}\beta^{-/-}$  mice (Flamant *et al.* 2002), there were increased numbers of pituitary thyrotrophs reflecting the enhanced TSH secretion in TR $\alpha 1^{-/-}\beta^{-/-}$  mice (Gothe *et al.* 1999).

The growth-promoting actions of GH have been shown to occur, in part, via IGF-I (Ohlsson *et al.* 1998). IGF-I and GH levels were reduced in TR $\alpha 1^{-/-}\beta^{-/-}$  mice and may be a cause of retarded growth (Gothe *et al.* 1999). The decrease observed in GH mRNA in TR $\alpha 1^{-/-}\beta^{-/-}$  mice, but not in TR $\alpha 1^{-/-}$  or TR $\beta^{-/-}$  single receptor knockout models suggests that both TR $\alpha 1$  and

**Table 2** Transgenic over-expression of mutated TR $\beta$ 

	Reference	Mutation	Promoter	Expression pattern	Thyroid status
<b>Model</b>					
G345R	Hayashi <i>et al.</i> (1998)	G345 (human)	TSH $\beta$	Pituitary	RTH
$\Delta$ 337T	Abel <i>et al.</i> (1999b)	$\Delta$ 337T (human)	Glycoprotein $\alpha$ -subunit	Pituitary	RTH
Cardiac $\Delta$ 337T	Pazos-Moura <i>et al.</i> (2000)	$\Delta$ 337T (human)	$\alpha$ -MHC	Cardiac	Euthyroid
PV mutant	Wong <i>et al.</i> (1997)	C insertion at nucleotide 1627	$\beta$ -actin	Ubiquitous	RTH
Pituitary PV mutant	Zhu <i>et al.</i> (1999)	C insertion at nucleotide 1627	Glycoprotein $\alpha$ -subunit	Pituitary	Euthyroid

MHC, myosin heavy chain.

TR $\beta$  have regulatory roles in GH gene transcription, although studies described above suggest that the effects of TR $\beta$  may predominate.

#### Transgenic models of human RTH

RTH has been classified as generalised (GRTH) or pituitary (PRTH) according to the absence or presence of thyrotoxicosis. Hayashi *et al.* (1998) selected the natural TR $\beta$  mutant G345R which causes severe RTH in man (Sakurai *et al.* 1989), and produced a transgenic mouse with the mutant TR targeted to the pituitary, under the control of the TSH $\beta$  promoter (Table 2). Pituitary-specific expression of TR $\beta$ <sup>G345R</sup> resulted in PRTH with increased serum T4 and reduced cholesterol levels but normal TSH concentrations. These findings reflected thyrotroph resistance to T3 and normal hepatic responsiveness, although the phenotype was milder than predicted from the human disease. TR $\beta$ <sup>G345R</sup> transgenic mice were normal in terms of fertility, survival and growth (Hayashi *et al.* 1998).

The naturally occurring mutation  $\Delta$ 337T (Usala *et al.* 1991, Baniahmad *et al.* 1992) was introduced into human TR $\beta$ 1 (Abel *et al.* 1999b). The TR $\beta$  <sup>$\Delta$ 337T</sup> mutant does not bind T3 and is associated with severe GRTH (Usala *et al.* 1991). Abel *et al.* (1999b) generated TR $\beta$  <sup>$\Delta$ 337T</sup> transgenic mice, selectively targeted to the pituitary via the common  $\alpha$ -subunit promoter (Table 2). There was a 3-fold increase in TSH but normal circulating T4 suggesting the possibility of reduced TSH bioactivity. Previous studies have indicated that TRH enhances the biological activity of TSH via alteration of TSH glycosylation (Beck-Peccoz *et al.* 1985, Taylor *et al.* 1988, Taylor & Weintraub 1989). This concept was supported recently by Yamada *et al.* (1997) who demonstrated that mice with a targeted ablation of TRH have T4 levels 50% lower than wild-type mice despite 2-fold elevated TSH concentrations. RTH patients have been observed to have increased bioactive TSH (Persani *et al.* 1994) and, thus, differences in TRH production may explain differences between human RTH patients and pituitary-specific TR $\beta$  <sup>$\Delta$ 337T</sup> transgenic mice

(Abel *et al.* 1999b). TR $\beta$  <sup>$\Delta$ 337T</sup> animals also show impaired ligand-independent activation of TSH $\beta$  gene expression *in vivo* and deficient TSH responsiveness to hypothyroidism. The results indicate that the pituitary-specific expression of TR $\beta$  <sup>$\Delta$ 337T</sup> leads to impaired T3-mediated repression, with defective T3-independent activation of TSH production, and that resistance at the level of both the pituitary and the hypothalamus is required for thyroid hormone elevation in RTH (Abel *et al.* 1999b).

A third RTH mutant, TR $\beta$ <sup>PV</sup>, was introduced into transgenic mice under the control of the human  $\beta$ -actin promoter to ensure ubiquitous expression (Table 2). Mice displayed features of RTH with elevated thyroid hormone levels and inappropriately normal TSH (Wong *et al.* 1997). Subsequently, Zhu *et al.* (1999) generated a transgenic mouse with the TR $\beta$ <sup>PV</sup> mutant targeted to the pituitary using the common  $\alpha$ -subunit promoter. Male TR $\beta$ <sup>PV</sup> mutant mice exhibited impaired weight gain but there were no alterations in T3, T4 or TSH compared with wild-type mice. Levels of GH and IGF-I were also unaffected, suggesting that impaired growth might be mediated by alternative factors (Zhu *et al.* 1999).

A major problem with the transgenic approach is that the mutant TR $\beta$  is introduced and expressed under the control of a heterologous promoter that might be ubiquitous or tissue specific but does not possess the regulatory features of the endogenous TR $\beta$  promoter. Furthermore, random integration of the transgene into host DNA may result in altered expression of other unknown potential modifier genes; the mutant receptor is introduced in the presence of both normal alleles of the native receptor; and transgene copy number may be important. These features suggest that a transgenic approach is unlikely to produce a faithful mouse model of RTH in which to understand physiological activities of TR isoforms in specific tissues.

#### Models of RTH using targeted mutagenesis

Kaneshige *et al.* (2000) addressed these issues by targeting the TR $\beta$ <sup>PV</sup> mutation to the endogenous mouse TR $\beta$  gene

**Table 3** Knock-in models using homologous recombination

	Reference	Mutation	Other information	Thyroid status
<b>Model</b>				
TR $\beta$ <sup>PV</sup>	Kaneshige <i>et al.</i> (2000)	C insertion at 1627 bp of mouse TR $\beta$ 1; frameshift and generation of 463 aa mutant protein	Cre/lox P system	Profound RTH
TR $\alpha$ 1 <sup>PV</sup>	Kaneshige <i>et al.</i> (2001)	C insertion at 1180 bp of mouse TR $\alpha$ 1; frameshift and generation of 409 aa mutant protein		RTH

PV, mutation derived from a patient with severe resistance to thyroid hormone (RTH).

by homologous recombination to generate a TR $\beta$ <sup>PV</sup> 'knock-in' mutant model of RTH (Table 3). The PV mutation was derived from a patient with severe RTH, who had elevated T4 and T3 levels, normal TSH, short stature, goitre and tachycardia (Parrilla *et al.* 1991). The mutation in exon 10 is a C insertion at codon 448 which produces a frameshift of the carboxy-terminal 14 amino acids (aa) of TR $\beta$ , resulting in severe impairment of T3 binding and transactivation activity by the mutant TR $\beta$ <sup>PV</sup> protein, which acts as a dominant negative TR *in vitro* (Kaneshige *et al.* 2000).

Heterozygous TR $\beta$ <sup>PV/+</sup> mice showed biochemical features of RTH that were typical of those in human RTH, whilst homozygous TR $\beta$ <sup>PV/PV</sup> mutants had severe RTH with disruption of the pituitary–thyroid feedback axis and impaired growth (Kaneshige *et al.* 2000). T3 concentrations were elevated 2- and 9-fold in TR $\beta$ <sup>PV/+</sup> and TR $\beta$ <sup>PV/PV</sup> mice respectively, and T4 levels were increased 2.5- and 15-fold in heterozygous and homozygous mutant mice. RTH was profound in TR $\beta$ <sup>PV/PV</sup> mice in which 412-fold increased TSH concentrations were seen. TR $\beta$ <sup>PV</sup> mice possessed thyroid glands of increased size with extensive hyperplasia seen in homozygous mutants. The PV mutation interfered with the activities of wild-type receptors on the expression of both  $\alpha$ -glycoprotein subunit and TSH $\beta$  genes and GH expression was repressed to 20% of wild-type values in TR $\beta$ <sup>PV/PV</sup> pituitaries. In heterozygous TR $\beta$ <sup>PV/+</sup> mice, there was a reduced GH mRNA response to elevated T3 levels (Kaneshige *et al.* 2000). Previous biochemical data have shown that TR $\beta$ <sup>PV</sup> mutant receptors form inactive heterodimers with wild-type TR $\beta$  or TR $\alpha$ 1, with TR $\beta$  having a higher interaction affinity than TR $\alpha$ 1 for TR $\beta$ <sup>PV</sup> (Zhu *et al.* 1996a). However, several mechanisms can be proposed to account for the dominant negative effect of mutant TR $\beta$ <sup>PV</sup>. Thus, the mild pituitary–thyroid abnormalities in TR $\beta$ <sup>PV/+</sup> mice might result from the formation of inactive or repressive TR $\beta$ <sup>PV</sup>-containing heterodimers, from indirect interference with the actions of one or both TR $\alpha$ 1 and TR $\beta$  by TR $\beta$ <sup>PV</sup>, from haplo-insufficiency of wild-type TR $\beta$  or a combination of such factors (Kaneshige *et al.* 2000).

Kaneshige *et al.* (2001) investigated this by generating mice with the PV mutation in TR $\alpha$ 1 (Table 3). TR $\alpha$ 1<sup>PV/+</sup> mice are dwarfs and have impaired fertility with reduced litter size and frequency of pregnancy. Although viable, mortality was high, a clear difference from the TR $\beta$ <sup>PV</sup> mice. TR $\alpha$ 1<sup>PV/PV</sup> homozygotes were especially rare after mating TR $\alpha$ 1<sup>PV/+</sup> animals and when born they died in the early neonatal period. TR $\alpha$ 1<sup>PV/+</sup> mice had normal T4, but elevated T3 and TSH levels with no histological abnormality of the thyroid gland. The elevated TSH, together with reduced T4:T3 ratio, suggests that mild thyroid failure occurs in TR $\alpha$ 1<sup>PV/+</sup> mice. There was no effect observed on pituitary GH expression but a 2- to 3-fold increased common  $\alpha$ -subunit mRNA expression was observed. TSH $\beta$  mRNA was unaffected, suggesting that increased circulating TSH concentrations may result from increased TSH protein translation, secretion or stability. The distinct phenotypes seen in the TR $\alpha$ 1<sup>PV</sup> (Kaneshige *et al.* 2001) and TR $\beta$ <sup>PV</sup> (Kaneshige *et al.* 2000) mice demonstrate that functions of the TR mutants are isoform specific *in vivo* and show that, although no TR $\alpha$  mutations have been documented in human RTH, dominant mutations affecting TR $\alpha$  are neither silent nor lethal. Thus, it seems probable that mutations of TR $\alpha$  are likely to exist in man, but that their resulting biochemical and general phenotypes would be predicted to be quite distinct from the RTH associated with TR $\beta$  mutations.

### Skeleton and growth

TR $\alpha$ <sup>0/0</sup> homozygous mice are proportionately growth retarded with reduced weight gain and preservation of a normal weight-to-length ratio, whereas heterozygotes grow normally (Gauthier *et al.* 2001). Growth retardation results from delayed endochondral bone formation. There is a failure of progression of hypertrophic chondrocyte differentiation in the epiphyseal growth plate, which is disorganised and immature. Bone mineralisation is impaired and trabecular bone density in the tibial metaphysis is reduced compared with wild-type mice

(Gauthier *et al.* 2001). These features are similar to those seen in hypothyroidism (Stevens *et al.* 2000), although in the basal state  $TR\alpha^{0/0}$  mice are biochemically euthyroid and have normal levels of pituitary GH (Gauthier *et al.* 2001), suggesting a defect in skeletal T3 responsiveness due to deletion of  $TR\alpha$ .

$TR\alpha^{-/-}$ -null mice are growth retarded with markedly delayed endochondral ossification (Fraichard *et al.* 1997), although these features are not manifest until the end of the second postnatal week. By the fifth postnatal week,  $TR\alpha^{-/-}$  mice lose up to 50% body weight and die, failing to survive beyond weaning. Skeletal analyses revealed normal morphology with significantly delayed ossification and mineralisation and a hypoplastic bone marrow. Histomorphometry showed reduced bone mass and increased areas of disorganised epiphyseal cartilage containing reduced numbers of hypertrophic chondrocytes (Fraichard *et al.* 1997, Gauthier *et al.* 1999). In contrast, deletion of  $TR\alpha 1^{-/-}$  (Wikstrom *et al.* 1998) does not result in growth abnormality and no skeletal abnormalities have been reported in  $TR\alpha 1^{-/-}$  mutants.

$TR\alpha 2^{-/-}$  mice have a slightly reduced growth rate after 5 weeks of age. Nevertheless, pituitary GH mRNA and protein levels are not different from wild-type animals, although serum IGF-I concentrations are diminished. Analysis of the skeleton revealed no evidence of altered longitudinal bone growth or change in growth plate width in  $TR\alpha 2^{-/-}$  mice. However,  $TR\alpha 2^{-/-}$  and  $TR\alpha 2^{+/-}$  heterozygotes showed evidence of impaired mineralisation in endochondral, but not intra-membranous bones, manifest by reduced bone mineral density and content (Salto *et al.* 2001). There was evidence of reduced tibial cortical bone in  $TR\alpha 2^{-/-}$  mice, suggesting late-onset growth retardation as these parameters may result from reduced periosteal bone growth in adults. A possible explanation for the adult onset growth retardation with reduced cortical bone in  $TR\alpha 2^{-/-}$  mice is that lack of  $TR\alpha 2$  or over-expression of  $TR\alpha 1$  results in late-onset IGF-I deficiency (Salto *et al.* 2001).

Complete deletion of  $TR\beta$  or  $TR\beta 2$  results in PRTH as indicated above, but does not cause growth retardation and no skeletal abnormalities have been reported in  $TR\beta^{-/-}$  or  $TR\beta 2^{-/-}$ -null mice (Abel *et al.* 1999a, Gauthier *et al.* 1999).

Growth retardation in  $TR\alpha^{0/0}\beta^{-/-}$  double mutant mice is more severe than that seen in  $TR\alpha^{0/0}$  single mutant animals despite the observation that  $TR\beta^{-/-}$  mice grow normally (Gauthier *et al.* 2001). These data suggest that some important growth-promoting actions may be performed by  $TR\beta$  in  $TR\alpha^{0/0}$  mutant mice. Nevertheless, GH production is also diminished in  $TR\alpha^{0/0}\beta^{-/-}$  double mutants relative to wild-type and  $TR\alpha^{0/0}$  mice, suggesting an additional cause of the profound growth retardation in  $TR\alpha^{0/0}\beta^{-/-}$  animals (Gauthier *et al.* 2001). The phenotype of delayed bone maturation and impaired ossification was seen in  $TR\alpha^{0/0}\beta^{-/-}$  mice

and is similar to that in  $TR\alpha^{0/0}$  mice (Gauthier *et al.* 2001). Comparison of  $TR\alpha^{0/0}\beta^{-/-}$  double mutants with hypothyroid  $Pax8^{-/-}$  animals indicated that more severe growth retardation and further delay in endochondral ossification occurs in  $Pax8^{-/-}$  mice, suggesting that unliganded apo-TRs also exert deleterious effects during bone formation in the complete absence of thyroid hormones (Flamant *et al.* 2002).

$TR\alpha^{-/-}\beta^{-/-}$  mice exhibit a similar skeletal phenotype to  $TR\alpha^{-/-}$  animals (Gauthier *et al.* 1999). There are no morphological alterations at birth whilst, similar to  $TR\alpha^{-/-}$  mice (Fraichard *et al.* 1997), double mutants undergo growth arrest and die at 5 weeks of age. Examination of the skeleton demonstrated delayed endochondral ossification with reduced mineralisation at 2 weeks of age, findings similar to those in  $TR\alpha^{-/-}$  mice (Gauthier *et al.* 1999). At this age, thyroid hormone concentrations in  $TR\alpha^{-/-}\beta^{-/-}$  mice did not differ from wild-type mice, indicating that skeletal abnormalities did not result from circulating hypothyroidism but were due to the  $TR\alpha^{-/-}\beta^{-/-}$  deletion (Gauthier *et al.* 1999).

$TR\alpha 1^{-/-}\beta^{-/-}$  mice (Gothe *et al.* 1999) are growth retarded from 3 weeks of age and by adulthood weigh 30% less than wild-type mice, with a disproportionate reduction in long bone length, delayed ossification and disorganisation of the growth plate. The bone defects resemble hypothyroidism, although in hypothyroidism the width of the tibial growth plate is reduced (Lewinson *et al.* 1989), whereas in  $TR\alpha 1^{-/-}\beta^{-/-}$  mice it is the reverse (Gothe *et al.* 1999). The skeletal abnormalities in  $TR\alpha 1^{-/-}\beta^{-/-}$  mice are associated with inhibition of the GH-IGF-I axis. Interestingly, growth retardation in  $TR\alpha 1^{-/-}\beta^{-/-}$  mice is reversed following GH replacement, but the growth plate ossification abnormalities are not rescued (Kindblom *et al.* 2001), indicating that TRs provide important actions to regulate the activity of the GH-IGF-I axis and also exert direct effects on the skeleton.

Transgenic  $TR\beta^{PV}$  mutant mice exhibited impaired weight gain compared with wild-type mice that was sexually dimorphic. Impairment of weight gain was evident in males expressing the transgene in all tissues (Wong *et al.* 1997) or in the pituitary gland only (Zhu *et al.* 1999). In females, growth retardation was less prominent and did not occur in animals with restricted expression of  $TR\beta^{PV}$  in the pituitary.

Reduced bone length and impaired weight gain was also seen in  $TR\beta^{PV}$  knock-in mice (Kaneshige *et al.* 2000), in which significantly shortened tibiae and femora were documented in both  $TR\beta^{PV/+}$  and  $TR\beta^{PV/PV}$  mice, with homozygotes more severely affected but no sexual dimorphism identified. Homozygous mutants also displayed a reduced growth spurt by up to 40% between 3 and 7 weeks of age. No differences were seen in  $TR\beta^{PV/+}$  heterozygotes. The same mutation of  $TR\alpha 1$  in  $TR\alpha 1^{PV/+}$  mice (Kaneshige *et al.* 2001) resulted in severe growth impairment shortly after birth with resultant dwarfism.

Four-week-old male TR $\alpha$ 1<sup>PV/+</sup> mice weighed 40% less and were 17% shorter than wild-type siblings, whilst females weighed 30% less than wild-type mice and were 15% shorter (Kaneshige *et al.* 2001).

### Skeletal muscle

Hypo- and hyperthyroidism in humans cause skeletal muscle weakness with fatigue and T3 plays an important role during skeletal muscle development (Gambke *et al.* 1983) by repressing or activating expression of MHC genes (Yu *et al.* 2000). Johansson *et al.* (2000) found that alterations in thyroid hormone levels induce changes in maximum shortening velocity (V0) and in isometric twitch contraction or relaxation times. V0 depends mainly upon myosin composition of the contractile proteins, the slow type MHC I and the faster MHC II isoforms (Schiaffino *et al.* 1989, Ennion *et al.* 1995). Thyrotoxicosis induces a shift towards faster MHC isoforms (Caiozzo *et al.* 1993) and hypothyroidism reverses this equilibrium (Johansson *et al.* 2000). Contraction and relaxation times also depend on the Ca<sup>2+</sup> transport capacity of the sarcoplasmic reticulum Ca<sup>2+</sup>-ATPase (SERCa), of which there are fast (SERCa1) and slow (SERCa2) types in skeletal muscle (Johansson *et al.* 2000). Hypothyroidism causes a decrease in both SERCa1 and SERCa2 activity (Sayen *et al.* 1992); this trend is reversed in thyrotoxicosis (Muller *et al.* 1994).

The actions of T3 in skeletal muscle have been studied in only a few of the TR mutant mice. TR $\alpha$ 1<sup>-/-</sup> mice exhibit a hypothyroid phenotype with longer muscle contraction and relaxation times and soleus muscle twitches extended by 40%. In TR $\beta$ <sup>-/-</sup> mice, the soleus muscle was less fatigue resistant than in wild-type mice although contraction and relaxation times were unchanged (Johansson *et al.* 2000). SERCa1 expression was reduced in TR $\alpha$ <sup>-/-</sup> mice but unaffected in TR $\beta$ <sup>-/-</sup> animals. In the soleus, total fibre number was reduced in TR $\alpha$ 1<sup>-/-</sup>  $\beta$ <sup>-/-</sup> mice: a 51% increase in MHC I fibres was seen in TR $\alpha$ 1<sup>-/-</sup>  $\beta$ <sup>-/-</sup> double mutant mice, a 21% difference was recorded in TR $\alpha$ 1<sup>-/-</sup> animals but no change was evident in TR $\beta$ <sup>-/-</sup> mice. These results corresponded with lower proportions of fast MHC II fibres in TR $\alpha$ 1<sup>-/-</sup>  $\beta$ <sup>-/-</sup> and TR $\alpha$ 1<sup>-/-</sup> mice (Yu *et al.* 2000) and suggest that both TR $\alpha$ 1 and TR $\beta$  are required for normal skeletal muscle responses to T3 (Johansson *et al.* 2000).

### Visual and auditory systems

Retinal cone photoreceptors facilitate colour vision (Ng *et al.* 2001). Cones expressing different opsin photopigments are sensitive to short (S, 'blue') and middle (M, 'green') wavelengths in rodents, and these cones are

differentially localised in the retina (Wang *et al.* 1992, Szel *et al.* 2000). Thyroid hormone has been implicated in the differentiation of cones (Kelley *et al.* 1995a,b) and mutations of TR $\beta$  may be associated with cone disorders in man (Newell & Diddie 1977). TR $\beta$ 1 is widely expressed, whereas the expression of TR $\beta$ 2 is largely restricted to the hypothalamus, pituitary, inner ear (Hodin *et al.* 1989, Bradley *et al.* 1994) and the outer nuclear layer of the embryonic retina that contains developing photoreceptors (Sjoberg *et al.* 1992). The mouse retina contains S- and M-opsin specific cones as well as cones expressing both (Rohlich *et al.* 1994, Szel *et al.* 1994, Lyubarsky *et al.* 1999). Deletion of TR $\beta$ 2 causes a selective loss of M-cones together with an increase in S-opsin immunoreactive cones. The cone distribution gradient is also disturbed with the usually focussed S-cones becoming widespread across the TR $\beta$ 2<sup>-/-</sup> retina. The presence of S-opsin in all TR $\beta$ 2<sup>-/-</sup> photoreceptors indicates that, in the absence of TR $\beta$ 2, all cones follow a default S-cone pathway and suggests a critical role for TR $\beta$ 2 in the commitment of photoreceptors to an M-cone or 'dual-cone' phenotype. The effect of TR $\beta$ 2 in cone photodevelopment is not compensated by TR $\beta$ 1 (Ng *et al.* 2001).

Thyroid hormones are also critical for development of the auditory system, an observation supported by clinical findings that demonstrate frequent associations between congenital thyroid disorders and hearing impairment or deafness (Refetoff *et al.* 1967, Stanbury 1984, DeLong 1993). Hypothyroidism has been reported to cause cochlear malformation (Deol 1973, Uziel *et al.* 1981). Recent studies reveal a period extending from late embryonic development to the second postnatal week where T3 is required for normal cochlea development (Deol 1973, Hebert *et al.* 1985, Uziel 1986). Exposure to thyroid hormone excess during the same period in rats causes an accelerated onset of auditory function development (Freeman *et al.* 1993). The cochlea appears to be the major T3-sensitive auditory organ and TR $\alpha$  mRNA is widely distributed in both cochlear and vestibular structures in the developing inner ear (Bradley *et al.* 1994). TR $\beta$  expression is more restricted and predominantly seen in the organ of Corti, which contains primordial hair cells, supporting cells and cells secreting tectorial membrane components. Both TR $\alpha$  and TR $\beta$  have been implicated in development of auditory function (Forrest *et al.* 1996a), although TR $\beta$  plays the major role.

TR $\beta$ <sup>-/-</sup> mice exhibit impaired auditory function. Auditory-evoked brainstem response (ABR) studies (Forrest *et al.* 1996a) have demonstrated that the threshold sound pressure levels required for detection of an ABR were significantly elevated for click stimuli and for all tones tested. Ninety-five percent of TR $\beta$ <sup>-/-</sup> mice tested were severely impaired with 10% of these being completely deaf and lacking a response to any stimulus. In mice with some preservation of hearing, the amplitude of ABR

waveforms was diminished although their pattern was normal, suggesting that the primary defect in  $TR\beta^{-/-}$  mice lies in the cochlea. The findings indicated that  $TR\beta$  controls the maturation of auditory function but is not required for cochlear morphogenesis (Forrest *et al.* 1996a). Recent data suggest that the basis for hearing impairment in  $TR\beta$ -null mice is a delay in the development of the expression of the fast-activating potassium conductance (IK,f) inner hair cells of the cochlea (Rusch *et al.* 1998). In wild-type mice, half-maximal IK,f expression is reached at postnatal day 17.5 whereas, in  $TR\beta^{-/-}$  animals, IK,f expression is absent up to day 18. This difference, at a critical developmental period, may explain the auditory phenotype of  $TR\beta$ -null mice (Rusch *et al.* 1998).  $TR\beta 1$  and  $TR\beta 2$  isoforms are coexpressed in the developing cochlea, but studies in  $TR\beta 2^{-/-}$  mice (Abel *et al.* 1999a) revealed no hearing deficit, indicating that  $TR\beta 2$  is not necessary for cochlear development and function.

Rusch *et al.* (1998) determined ABR thresholds in  $TR\alpha 1^{-/-}$  knockout mice (Wikstrom *et al.* 1998). Over a wide range of frequency stimuli, these mice show no impairment, indicating that  $TR\alpha 1$  is not essential for auditory function (Rusch *et al.* 1998).  $TR\alpha^{0/0}$  mice exhibit a normal hearing threshold at low or middle frequencies, but a marked sensitivity loss at high tone frequencies (Gauthier *et al.* 2001). In  $TR\alpha^{0/0}\beta^{-/-}$  mice, the effects on hearing thresholds were significantly more severe than in  $TR\beta^{-/-}$  mice (Gauthier *et al.* 2001). In summary,  $TR\beta 1$  appears to be the major mediator of T3 action in the ear, although  $TR\alpha$  is also required for full auditory function.

### Small intestine

Thyroid hormones influence postnatal intestinal maturation by stimulating intestinal crypt cell proliferation and the onset of brush border enzyme expression (Castillo *et al.* 1991, Hodin *et al.* 1996). Thus, T3 enhances the dramatic developmental changes in the gastrointestinal tract that lead to its complete remodelling by adulthood (Tata 1993, Shi *et al.* 1996).

The  $TR\Delta\alpha 1$  and  $\Delta\alpha 2$  mRNA isoforms are transcribed from an internal promoter, located in intron 7 of the *TRA* gene (Chassande *et al.* 1997). Their expression is highest in a limited number of organs including the lung and small intestine (Fraichard *et al.* 1997).  $TR\Delta\alpha 1$  mRNA is expressed mainly in the epithelium and lamina propria of the distal ileum, with barely detectable levels being seen in the muscular layers of the distal ileum and proximal colon.  $TR\Delta\alpha 2$  mRNA is similarly distributed, except that maximal expression occurs in the lamina propria of the proximal jejunum (Plateroti *et al.* 2001).

Fraichard *et al.* (1997) described delayed small intestine development in  $TR\alpha^{-/-}$  mice with no overall alterations in stomach, duodenum or colon. The small intestine

was smaller, softer and more fragile than in wild-type mice. The diameters of the jejunum and ileum were reduced, together with the number and size of villi, and the total number of epithelial cells per crypt villus unit was reduced by 65% in the ileum. The frequency of goblet cells was reduced 2.5-fold in the ileum of  $TR\alpha^{-/-}$  animals, but Paneth cell frequency was unaffected. Circular and smooth muscle layers were reduced, correlating with altered motility in the  $TR\alpha^{-/-}$  ileum, which exhibited no spontaneous contraction or response to pharmacological stimuli. Small intestine dipeptidase and disaccharidase activities were reduced 2-fold in  $TR\alpha^{-/-}$  mice relative to wild-type mice, although peptide and disaccharide absorption was normal. These are features of delayed intestinal maturation as a similar phenotype is characteristic in young mice.

The small intestine was further characterised by Plateroti *et al.* (1999). In  $TR\alpha^{-/-}$  mice, the crypt-villus length was shortened in proximal jejunum and distal ileum compared with wild-type mice. These findings correlate with a decreased proliferation rate of  $TR\alpha^{-/-}$  intestinal crypt cells and a decreased number of goblet cells in the  $TR\alpha^{-/-}$  ileum. Furthermore,  $TR\alpha^{-/-}$  villous epithelial cells had reduced cellular and brush border height compared with wild-type mice. mRNA expression of the intestine-specific Cdx-1 and Cdx-2 homeobox genes was reduced in the  $TR\alpha^{-/-}$  mice. These transcription factors are expressed normally in a specific pattern with an increasing gradient along the longitudinal axis of the gut (James & Kazenwadel 1991, Freund *et al.* 1992). Cdx-1 and Cdx-2 play a key regulatory role in normal intestinal cell proliferation and differentiation (Traber 1994, Suh & Traber 1996, Duluc *et al.* 1997, Lorentz *et al.* 1997) and reduced expression in  $TR\alpha^{-/-}$  mice may account, in part, for the severe small intestine phenotype.

Plateroti *et al.* (1999) observed that a single T3 injection stimulated recovery of intestinal morphology in  $TR\alpha^{-/-}$  mice. Furthermore, digestive enzyme activities were partially rescued and Cdx-1 mRNA was restored to normal wild-type levels. Cdx-2 mRNA expression, however, was reduced. In follow-up studies, Plateroti *et al.* (2001) investigated mechanisms responsible for the intestinal abnormalities in  $TR\alpha^{-/-}$  mice. They retain expression of  $TR\Delta\alpha 1$  and  $\Delta\alpha 2$ , suggesting that expression of these isoforms in the absence of  $TR\alpha 1$  may be deleterious to intestinal development. Gauthier *et al.* (2001) found that, in  $TR\alpha^{0/0}$  animals, the size of the small intestine mucosa was reduced compared with wild-type animals, due to decreased villus length and reduced number of epithelial cells per crypt-villus axis. The features were not as severe as in  $TR\alpha^{-/-}$  mice (Plateroti *et al.* 2001). Furthermore, in  $TR\alpha^{0/0}$  mice, expression of Cdx-1 and Cdx-2 mRNAs remained unchanged relative to wild-type mice, indicating that the  $TR\Delta\alpha 1$  and  $\Delta\alpha 2$  mRNAs may exert an inhibitory role to regulate Cdx gene transcription in the absence of  $TR\alpha 1$ . Thus, the relative level of expression of

TR $\alpha$  and  $\Delta\alpha$  isoforms in the intestine may be a critical determinant in normal small intestinal development (Plateroti *et al.* 2001). In order to investigate this, TR $\alpha^{7/7}$  mutant mice were generated, in which expression of TR $\Delta\alpha 1$  and  $\Delta\alpha 2$  transcripts was almost completely abolished whilst expression of TR $\alpha 1$  and  $\alpha 2$  was retained at normal levels. These mice display no morphological alterations in the ileum compared with wild-type mice and, in contrast to both TR $\alpha^{-/-}$  and TR $\alpha^{0/0}$  mice, TR $\alpha^{7/7}$  mutants had increased numbers of goblet cells compared with wild-type mice, whilst levels of Cdx-1 and Cdx-2 mRNAs were unchanged (Plateroti *et al.* 2001). Thus, the severity of the intestinal phenotype correlates with the level of expression of TR $\Delta\alpha$  transcripts and not with the absence of TR $\alpha 1$  or TR $\alpha 2$  isoforms (Gauthier *et al.* 2001). Balanced expression of all TR $\alpha$  isoforms is essential for normal small intestine development at weaning (Plateroti *et al.* 2001).

TR $\beta^{-/-}$  mice display no intestinal phenotype, indicating that only TR $\alpha$  is essential for postnatal small intestine development (Plateroti *et al.* 1999). Indeed, ablation of TR $\beta$  in TR $\alpha^{0/0}\beta^{-/-}$  mice does not worsen the intestinal phenotype seen in TR $\alpha^{0/0}$  animals (Gauthier *et al.* 2001). In contrast, however, TR $\alpha^{-/-}\beta^{-/-}$  mice have more severe intestinal abnormalities than their TR $\alpha^{-/-}$  counterparts (Gauthier *et al.* 1999, Plateroti *et al.* 1999). Despite high circulating thyroid hormone levels, there was a marked decrease in the number of epithelial and goblet cells per crypt-villus unit in the distal ileum of TR $\alpha^{-/-}\beta^{-/-}$  mice, together with severely reduced expression of lactase mRNA and protein and Cdx-1 and Cdx-2 mRNAs (Plateroti *et al.* 1999). These data demonstrate that ablation of the TR $\beta$  gene significantly worsens the phenotype observed in TR $\alpha^{-/-}$  mice. Thus, expression of the non-T3-binding TR $\Delta\alpha$  variants in the absence of all T3-binding TR $\alpha$  and  $\beta$  isoforms is deleterious to intestinal development, confirming further that a balance between hormone-bound TRs and apo-TR $\Delta\alpha$  isoforms is essential for intestinal development. Interestingly, distal small intestine epithelial cell proliferation and differentiation are more compromised in Pax8 $^{-/-}$  mice relative to TR $\alpha^{0/0}\beta^{-/-}$  double mutants, although the phenotype can be completely rescued in Pax8 $^{-/-}$  mice following T3 administration. Furthermore, in compound Pax8 $^{-/-}$ TR $\beta^{-/-}$  mutants, a severe phenotype similar to that in Pax8 $^{-/-}$  mice is seen, whereas in Pax8 $^{-/-}$ TR $\alpha^{0/0}$  compound mutants there is partial rescue (Flamant *et al.* 2002). These findings provide evidence to support the hypothesis that the TR $\Delta\alpha$  isoforms play the key role in postnatal small intestinal development (Flamant *et al.* 2002).

## Liver

TR $\beta$  is the major functional TR in liver, accounting for 80% of hepatic T3-binding activity, with the remainder

resulting from TR $\alpha 1$  (Weiss *et al.* 1998). Accordingly, TR $\beta^{-/-}$  mice display altered hepatic T3 responsiveness (Weiss *et al.* 1998). Thyroid hormone deprivation causes reduced expression of spot 14 (S14) in wild-type mice, but no change is seen in TR $\beta^{-/-}$  mice. Similarly, T3 stimulates S14 and malic enzyme (ME) expression in wild-type animals but not in TR $\beta^{-/-}$  mice. One of the best characterised hepatic T3 target genes is the type I 5'-deiodinase enzyme (D1) which catalyses the conversion of T4 to T3 in liver and kidney (Sato *et al.* 1984, Maia *et al.* 1995). D1 is induced by T3 and the human promoter contains several well-studied T3 response elements (Toyoda *et al.* 1995). Amma *et al.* (2001) studied D1 expression in TR-deficient mice and showed, in TR $\beta^{-/-}$  mice, that liver and kidney D1 mRNA levels were reduced to less than 30% and 50% of wild-type levels respectively. Thus, TR $\beta$  is required for the maintenance of basal D1 expression. The finding that liver D1 mRNA is undetectable in TR $\alpha 1^{-/-}\beta^{-/-}$  mice, with enzyme activity less than 0.03% of wild-type mice (Amma *et al.* 2001), suggests, however, that TR $\alpha$  and TR $\beta$  exert some common functions in liver, although TR $\beta$  appears predominant. In accord with this, TR $\alpha 1^{-/-}$  mice were found to have a 2-fold elevation in liver D1 expression, presumably resulting from increased TR $\beta$  activity due to elevated T3 concentrations in these animals. Interestingly, renal D1 mRNA expression was not elevated in TR $\alpha 1^{-/-}$  mice (Wikstrom *et al.* 1998), suggesting that the kidney is predominantly a TR $\alpha$ -sensitive target organ.

Flamant *et al.* (2002) reported that hepatic D1 expression is significantly reduced in TR $\alpha^{0/0}\beta^{-/-}$  mice and undetectable in Pax8 $^{-/-}$  mutants. The presence of D1 expression in the TR $\alpha^{0/0}\beta^{-/-}$  liver but absent expression in Pax8 $^{-/-}$  mutants indicates that unliganded TRs may actively repress basal transcription of D1. Both Pax8 $^{-/-}$ TR $\alpha^{0/0}$  and Pax8 $^{-/-}$ TR $\beta^{-/-}$  compound mutants do not express detectable D1, implicating both TR $\alpha$  and TR $\beta$  unliganded receptors in active repression of the D1 promoter.

Expression of the mutant TR $\beta^{PV}$  which results in RTH was also observed to suppress the magnitude of hepatic T3-target gene responses. Expression of S14, ME and D1 was significantly reduced in TR $\beta^{PV/+}$  heterozygotes compared with wild-type levels and a further reduction in ME and S14, together with undetectable D1 expression, was evident in TR $\beta^{PV/PV}$  mutants (Kaneshige *et al.* 2000). These findings contrast with results obtained in TR $\beta$ -null mice, where no change in ME expression was detected (Weiss *et al.* 1998). In contrast, a different profile of abnormal T3 target gene regulation was reported in TR $\alpha 1^{PV/+}$  mice. Activity of ME and D1 was increased 2.2-fold and 9.2-fold respectively, indicating hypersensitivity to T3 (Kaneshige *et al.* 2001). These findings may result from the markedly increased T3 concentrations in TR $\alpha 1^{PV/+}$  mice, which may overcome some of the dominant negative activity of the  $\alpha 1^{PV}$  mutant receptor

and increase T3 responses mediated via TR $\beta$  (Kaneshige *et al.* 2001).

Hepatic cholesterol metabolism is also influenced by T3, which regulates cholesterol 7 $\alpha$ -hydroxylase (CYP7A), the rate-limiting enzyme in bile acid synthesis (Gullberg *et al.* 2000). Hypothyroidism is associated with increased serum cholesterol, especially affecting low density lipoprotein (LDL) cholesterol, which can be normalised following T4 replacement (O'Brien *et al.* 1993). T3 also influences hepatic synthesis and uptake of cholesterol, controlled largely by hepatic 3-hydroxy-3-methylglutaryl coenzyme A reductase and LDL receptors (Ness *et al.* 1973, 1990, Mathe & Chevallier 1976, Staels *et al.* 1990, Ness & Zhao 1994), and the degradation of cholesterol into bile acids, where the rate-limiting CYP7A is transcriptionally induced by T3 (Mathe & Chevallier 1976, Crestani *et al.* 1994, Pandak *et al.* 1997). The normal stimulation of CYP7A mRNA and enzyme activity is unaffected in TR $\alpha$ 1<sup>-/-</sup> mice, whereas T3 responsiveness is absent in TR $\beta$ <sup>-/-</sup> animals, indicating that TR $\beta$  plays a key role in the regulation of hepatic cholesterol metabolism. Further work, however, has demonstrated an enhanced CYP7A response in T3-deficient TR $\beta$ <sup>-/-</sup> mice challenged with dietary cholesterol such that mutant mice did not develop hypercholesterolaemia to the same extent as wild-type controls. The results suggest that TRs influence a range of regulatory effects that may be independent of circulating T3 (Gullberg *et al.* 2000).

### Heart and thermogenesis

Thyroid hormones influence cardiac conductance, heart rate, cardiac output and cardiac muscle growth. Hypothyroidism leads to bradycardia, whereas hyperthyroidism induces tachycardia. Thyroid hormone plays a crucial role in the control of thermogenesis with hypothyroidism leading to reduced oxygen consumption and heat production. The action of T3 in the heart is mediated predominantly by TR $\alpha$ 1 (Gloss *et al.* 2001) and TR $\beta$ 1 is expressed in the heart at low levels (Falcone *et al.* 1992). Wikstrom *et al.* (1998) reported that TR $\alpha$ 1-deficient mice have a lower heart rate than control animals of the same genetic background. However, TR $\alpha$ 1<sup>-/-</sup> mice are hypothyroid and when hyperthyroidism is induced the heart rate increases in both TR $\alpha$ 1<sup>-/-</sup> and wild-type animals. Nevertheless, the response in TR $\alpha$ 1<sup>-/-</sup> mice is sub-optimal indicating that ablation of TR $\alpha$ 1 reduces intrinsic heart rate and cardiac responsiveness to T3. Thus, TR $\beta$  may influence heart rate via mechanisms that are distinct from the actions of TR $\alpha$ 1 or that compensate for it. Electrocardiograph (ECG) recordings revealed a prolonged QRS complex and QT end duration in TR $\alpha$ 1<sup>-/-</sup> mice, findings that are similar to those in hypothyroid rat cardiomyocytes (Shimoni & Severson 1995) and hypothyroid patients.

Thyroid hormones control body temperature by stimulating gluconeogenesis and increasing thermogenesis in brown adipose tissue. The 24-hour mean core body temperature was 0.5 °C lower in TR $\alpha$ 1<sup>-/-</sup> mice, with no significant difference in locomotor activity compared with heterozygotes and wild-type animals. TR $\alpha$ 1<sup>-/-</sup> mice also have normal brown adipose tissue mass, suggesting that reduced body temperature results from altered metabolic activity (Wikstrom *et al.* 1998). TR $\beta$ <sup>-/-</sup> mice, however, have normal body temperature suggesting that the involvement of TR $\beta$  in basal thermogenesis is minor, or may be substituted by TR $\alpha$ 1 (Gauthier *et al.* 2001). There is also a slight increase in heart rate in TR $\beta$ <sup>-/-</sup> mice compared with wild-type mice that is not relieved by T3 administration (Weiss *et al.* 1998, Johansson *et al.* 1999). TR $\beta$ <sup>-/-</sup> mice have a reduced QT end time, but exhibit no other ECG abnormalities (Johansson *et al.* 1999). HCN2 and HCN4 are cardiac genes that contribute to pacemaker activity by coding for the hyperpolarisation-activated current (Ludwig *et al.* 1998). In TR $\beta$ <sup>-/-</sup> mice, levels of HCN2 and HCN4 are significantly increased, leading to an increase in contractile function. However, when TR $\beta$ <sup>-/-</sup> mice were rendered euthyroid, heart rate and HCN gene expression was normalised (Gloss *et al.* 2001), suggesting that T3 responsiveness is mediated via TR $\alpha$ . Furthermore, in Pax8<sup>-/-</sup> mice, HCN2 mRNA expression is extremely low, although it is inducible by T3. In compound Pax8<sup>-/-</sup> TR $\alpha$ 0<sup>0/0</sup> mutants, HCN2 expression is restored whereas in Pax8<sup>-/-</sup> TR $\beta$ <sup>-/-</sup> animals expression remains low. These results indicate that repression of basal HCN2 expression is mediated via unliganded TR $\alpha$ 1 (Flamant *et al.* 2002).

Deletion of both TR $\alpha$ 1 and TR $\alpha$ 2 in TR $\alpha$ <sup>-/-</sup> mice results in bradycardia and decreased HCN2 and HCN4 gene expression, together with reduced myocardial contractility (Gloss *et al.* 2001). TR $\alpha$ 0<sup>0/0</sup> mice, similarly, exhibit a reduced basal heart rate (Macchia *et al.* 2001) and also have a 0.4 °C reduction in body temperature (Gauthier *et al.* 2001). Reduced cardiac sensitivity to dynamic testing in TR $\alpha$ 0<sup>0/0</sup> mice indicates that TR $\alpha$  is the major functional cardiac TR isoform (Macchia *et al.* 2001). Furthermore, body temperature data from TR $\alpha$ 0<sup>0/0</sup> mice imply that TR $\alpha$ 2 has no effect on thermogenesis under basal conditions (Gauthier *et al.* 2001). TR $\alpha$ 1<sup>-/-</sup>  $\beta$ <sup>-/-</sup> animals have a similar phenotype to TR $\alpha$ <sup>-/-</sup> animals, exhibiting basal bradycardia, prolonged PQ and QT end durations, and a 0.4 °C reduction in body temperature relative to wild-type mice (Johansson *et al.* 1998, 1999). TR $\alpha$ 0<sup>0/0</sup>  $\beta$ <sup>-/-</sup> mice have a marked mean body temperature reduction of 4 °C compared with wild-type mice (Gauthier *et al.* 2001) and reduced HCN2 gene expression (Flamant *et al.* 2002). Thus, comparison of temperature reductions in TR $\alpha$ <sup>-/-</sup>  $\beta$ <sup>-/-</sup> (0.5 °C) and TR $\alpha$ 0<sup>0/0</sup>  $\beta$ <sup>-/-</sup> (4 °C) mice reveals that TR $\alpha$ 2 may exert important actions to regulate thermogenesis (Gauthier *et al.*

2001). Nevertheless, in  $TR\alpha 2^{-/-}$  mice (Salto *et al.* 2001) a 10% decrease in basal heart rate together with a 0.4 °C increase in body temperature was reported. These data are difficult to interpret, however, because of the overexpression of  $TR\alpha 1$  in  $TR\alpha 2^{-/-}$  mice that results from the targeting strategy.

In  $TR\beta^{\Delta 337T}$  transgenic mice, two groups assessed the cardiac phenotype by expressing the transgene either generally (Gloss *et al.* 1999) or specifically in the myocardium (Pazos-Moura *et al.* 2000). The expression of cardiac myosin genes is influenced by T3 (Hoh *et al.* 1978, Everett *et al.* 1984, Klein & Hong 1986). In euthyroidism,  $\alpha$ MHC is the predominant MHC isoform, its expression is increased in thyrotoxicosis, whereas  $\beta$ MHC is predominant in hypothyroidism (Izumo *et al.* 1986). In both  $\Delta 337T$  transgenic models, basal heart rate was reduced. Expression of  $\beta$ MHC was increased together with a parallel reduction in  $\alpha$ MHC expression and ECG studies demonstrated broadening of the QRS complex, indicating prolongation of ventricular depolarisation time. In myocardium-specific  $\Delta 337T$  mice, the PQ, QR and ST intervals were also prolonged (Pazos-Moura *et al.* 2000), which was not the case in mice harbouring the ubiquitously expressed  $\Delta 337T$  transgene (Gloss *et al.* 1999); although both models displayed a hypothyroid pattern of gene expression, ECG changes and contractile function despite normal T3, T4 and TSH levels.

### Central nervous system (CNS) and behaviour

The importance of thyroid hormones in the development of the CNS is well known. TRs are distributed widely throughout the brain, in neurons and glial cells, and the profile of TR expression in different brain regions has been suggested to implicate specific functions for different TR isoforms in CNS development (Forrest *et al.* 1990, Bradley *et al.* 1992).

In  $TR\alpha^{-/-}$  mice, brain size was marginally reduced, but no morphological or cellular differences were observed (Fraichard *et al.* 1997). No differences were noted in laminar organisation or brain cytoarchitecture by Nissl staining. Histochemical analysis of acetylcholine esterase and cytochrome oxidase showed no differences in cortical structure or subcortical nuclear organisation between  $TR\alpha^{-/-}$  and wild-type mice (Fraichard *et al.* 1997). Studies were also performed in  $TR\beta^{-/-}$  mice (Forrest *et al.* 1996b).  $TR\beta^{-/-}$  animals completed the Morris water maze test with equal proficiency to wild-type controls. Open field and Y-Maze testing produced similar results. No abnormalities in brain anatomy, including T3-sensitive structures such as the cerebellum and hippocampus, were observed following histological analysis. No overt neurological defects were seen, indicating that the  $TR\beta$  gene may have a subtle rather than a major role in CNS development (Forrest *et al.* 1996b).  $TR\beta$  is induced

in neonatal CNS at a critical T3-sensitive period (Forrest *et al.* 1991, Mellstrom *et al.* 1991) and the rise in  $TR\beta 1$  occurs with the onset of the T3-dependent regulation of genes including myelin basic protein (MBP) and Purkinje cell protein-2 (Pcp-2) (Strait *et al.* 1992). However, whilst the onset of MBP expression is delayed in the hypothyroid mouse (Farsetti *et al.* 1992), no differences in MBP or Pcp-2 mRNA levels were observed in  $TR\beta^{-/-}$  mice compared with wild-type controls, suggesting that  $TR\beta$  is not critical for the normal postnatal expression of these T3-dependent genes (Sandhofer *et al.* 1998). Taken together, these data suggest that  $TR\alpha$  and  $TR\beta$  mediate redundant actions in the CNS.

To investigate further,  $TR\alpha 1^{-/-}\beta^{-/-}$  mice, lacking both  $TR\alpha 1$  and  $TR\beta$ , were examined (Gothe *et al.* 1999). Although these mice show hypothyroid features, they have normal activity and do not fatigue (Johansson *et al.* 1999). Macroscopic examination of  $TR\alpha 1^{-/-}\beta^{-/-}$  brains showed no gross cellular or morphological abnormalities compared with wild-type controls. However, TRH mRNA levels were markedly elevated in the paraventricular nucleus (PVN) and medullary raphe nuclei of  $TR\alpha 1^{-/-}\beta^{-/-}$  animals, whereas substance P mRNA, a peptide coexpressed with TRH in the medullary raphe nuclei, was unchanged. Differential regulation of TRH receptor mRNA was observed in specific regions of the  $TR\alpha 1^{-/-}\beta^{-/-}$  brain, with levels increased in the motor neurons of the trigeminal nucleus, unchanged in subiculum and decreased in the amygdala. Galanin mRNA was also reduced in the PVN of  $TR\alpha 1^{-/-}\beta^{-/-}$  mice, although levels were unchanged in the dorsomedial and arcuate nuclei. In the arcuate nucleus, glutamic acid decarboxylase (GAD) and prepro-neuropeptide Y (NPY) mRNA levels were not altered. However, tyrosine hydroxylase (th) mRNA levels were decreased. In the olfactory bulb, reduced calbindin immunostaining in the glomerular layer with a decrease of calbindin-immunoreactive neurons was seen in  $TR\alpha 1^{-/-}\beta^{-/-}$  mice, and reduced mRNA levels were recorded in the glomerular layer. Also, GAD mRNA levels were selectively reduced in the glomerular layer of the olfactory bulb of double mutant mice, but galanin and NPY mRNA expression was unchanged. The regulation of TRH signalling pathways in  $TR\alpha 1^{-/-}\beta^{-/-}$  mice is, therefore, complex and regionally selective. In  $TR\alpha 1^{-/-}\beta^{-/-}$  animals, the reduction in calbindin- and th-positive dendrites may also indicate impaired neuronal maturation in the olfactory bulb (Calza *et al.* 2000). Galanin expression is also selectively altered in the PVN of double mutant mice. It has a role in the regulation of pulsatile GH secretion (Maiter *et al.* 1990) and reduced galanin mRNA levels in the PVN may be involved in growth impairment seen in  $TR\alpha 1^{-/-}\beta^{-/-}$  mice (Calza *et al.* 2000). Since thyroid hormones are such key regulators of CNS development, it is surprising that more extensive abnormalities were not observed in  $TR\alpha 1^{-/-}\beta^{-/-}$  mice. It is likely that

redundancy between complex compensatory pathways may account for the minimal phenotype.

In behavioural tests,  $TR\beta^{-/-}$  animals show no abnormalities and no gross CNS phenotype is observed in  $TR\alpha$ - or  $TR\beta$ -deficient animals. However, recent data suggest that the  $TR\alpha$ 1 and  $TR\beta$  genes may have opposite effects on the sexual behaviour of female mice (Dellovade *et al.* 2000). Both molecular (Zhu *et al.* 1996b, Dellovade *et al.* 1999a,b) and behavioural (Dellovade *et al.* 1996) studies have shown that thyroid hormones affect the neural systems mediating oestrogen-dependent sexual behaviour in rodents. Dellovade *et al.* (2000) reported that  $TR\beta^{-/-}$  female mice show increased sexual behaviour, whereas, female  $TR\alpha$ 1 $^{-/-}$  animals display a reduced lordosis pattern. In the brain, the number of oestrogen-receptor immunoreactive neurons were similar across wild-type,  $TR\alpha$ 1 $^{-/-}$  and  $TR\beta^{-/-}$  genotypes. However, the number of oxytocin immunoreactive neurons in the PVN was increased in  $TR\beta^{-/-}$  animals, and a concomitant reduction was observed in  $TR\alpha$ 1 $^{-/-}$  mice (Dellovade *et al.* 2000). Oxytocin has been reported to facilitate a variety of reproductive and affiliative behaviours (Pedersen *et al.* 1992). Thus, oestrogen-influenced oxytocin immunoreactivity in the PVN may explain, in part, the behavioural results of  $TR\alpha$ 1 $^{-/-}$  and  $TR\beta^{-/-}$  mice. Although, in many physiological systems, there is co-operation between TRs, this specific behaviour pattern reveals functional antagonism between  $TR\alpha$ 1 and  $TR\beta$  (Dellovade *et al.* 2000).

In mice with the RTH  $TR\beta^{\Delta 337T}$  mutation, marked impairment in balance and co-ordination was observed (Hashimoto *et al.* 2001) following beam and rotorod testing (Scherbel *et al.* 1999). In the Morris water maze test (Forrest *et al.* 1996b), both heterozygous and homozygous mutants exhibited a learning deficit (Hashimoto *et al.* 2001). The defect was similar to that seen in congenital hypothyroidism (Anthony *et al.* 1993).  $TR\beta^{\Delta 337T}$  mice have a smaller cerebellum, although they exhibit no reduction in the size of the forebrain. Haematoxylin and eosin staining of the hippocampus and dentate gyrus revealed no gross morphological differences. However, hippocampal brain-derived neurotrophic factor immunoreactivity was reduced in heterozygous and homozygous mutants. The number of stained granule cells and fibres was especially reduced in the CA3 hippocampal region.  $TR\beta^{\Delta 337T}$  mice exhibit hypothyroid CNS changes, despite elevated thyroid hormone levels (Hashimoto *et al.* 2001), indicating a dominant negative effect of the mutant receptor *in vivo*. A transgenic mouse with  $TR\beta^{\Delta 337T}$  selectively targeted to the pituitary has also been developed (Abel *et al.* 1999b). These mice show significant down-regulation of prepro-TRH mRNA in the caudal paraventricular hypothalamus. Regional differences in TRH gene expression within the PVN and reduced PVN prepro-TRH mRNA were also observed, suggesting that hypothalamic TRH neurons are

extremely sensitive to changes in thyroid status (Abel *et al.* 1999b).

Transgenic mice expressing the PV mutant  $TR\beta$  exhibit a hyperactivity behavioural phenotype (Wong *et al.* 1997). A clinical feature of human RTH is attention deficit hyperactivity disorder, which affects 73% of children and 42% of adults (Brucker-Davis *et al.* 1995). The development of structure and function in the mammalian brain is accompanied by marked increases in local rates of glucose utilisation (Duffy *et al.* 1982, Kennedy *et al.* 1982), which may be sensitive to circulating thyroid hormone levels (Dow-Edwards *et al.* 1986). In targeted knock-in mutant mice,  $TR\beta^{PV}$  mRNA has been detected in cerebrum, cerebellum, pituitary gland and systemic organs (Kaneshige *et al.* 2000) and expression of  $TR\alpha$ 1 $^{PV}$  mRNA has been identified in both cerebrum and cerebellum (Itoh *et al.* 2001). Following determination by the quantitative autoradiographic 2-[ $^{14}C$ ]deoxyglucose method (Sokoloff *et al.* 1977), cerebral glucose utilisation was significantly lower in 4-week-old  $TR\alpha$ 1 $^{PV/+}$  mice compared with wild-type mice.  $TR\beta^{PV/PV}$  animals, of the same age, showed no change (Itoh *et al.* 2001). Expression of *Srg1*, a member of a protein family involved in the regulation of neurotransmitter release (Fernandez-Chacon *et al.* 2001) and a T3-positively regulated gene expressed in several brain regions (Thompson 1996), was reduced to 40% in the cerebellum of  $TR\alpha$ 1 $^{PV/+}$  animals, with no change seen in  $TR\beta^{PV/+}$  or  $TR\beta^{PV/PV}$  mice (Itoh *et al.* 2001). Abnormal expression of T3 target genes was also observed in the cerebellum of  $TR\alpha$ 1 $^{PV/+}$  mice. No change of MBP and *Pcp-2* expression was seen in  $TR\beta^{PV/+}$  animals, although activations of 1.4- and 1.2-fold were recorded for MBP and *Pcp-2* genes respectively in  $TR\alpha$ 1 $^{PV/+}$  mice (Kaneshige *et al.* 2001).

## Immune system

A role for thyroid hormones in B and T lymphocyte development has been revealed in Snell dwarf mice (Fabris *et al.* 1971, Murphy *et al.* 1992, 1993, Montecino-Rodriguez *et al.* 1996, 1997). Arpin *et al.* (2000) studied  $TR\alpha^{-/-}$  mice to determine T3 targets in the immune system and the TRs involved in lymphoid development. They found that the size of the mature B cell pool is altered due to a reduction in progenitor cell proliferation. In  $TR\alpha^{-/-}$  mice, no change in primary (thymus and bone marrow) or secondary (spleen) lymphoid organ cell numbers were seen at 10 days of age; however, by postnatal day 24, cell numbers in all lymphoid organs were reduced. The cellularity of T and B cells was reduced by 30% and 60% respectively, and the number of spleen granulocytes and macrophages was also reduced. There was also evidence of haemopoiesis in the  $TR\alpha^{-/-}$  spleen (Arpin *et al.* 2000).  $TR\alpha^{-/-}$  mice suffer from stress-induced defects, which may influence immune function.

Glucocorticoids down-regulate numerous immune processes and induce lymphocyte apoptosis (Wilder 1995). Cortisol levels were, therefore, measured in TR $\alpha^{-/-}$  mice, but no differences between TR $\alpha^{-/-}$  and wild-type mice were observed. Double positive thymocytes are sensitive to glucocorticoid-induced depletion of lymphocytes (Gruber *et al.* 1994) providing an objective index of stress. Up to 24 days of age, CD4<sup>+</sup>CD8<sup>+</sup> cells remained unaffected in TR $\alpha^{-/-}$  mice suggesting an absence of stress-induced immune modifications. Thus, T3 acts directly on B cells and this may be via control of progenitor cell proliferation or differentiation by TR $\alpha$ 1 or  $\alpha$ 2 (Arpin *et al.* 2000). Nevertheless, it has been proposed that T3 may act via stimulation of GH-IGF-I secretion from B cells (Weigent *et al.* 1992, Geffner 1997), but this appears unlikely since GH cannot restore the pre-B cell compartment in Snell mice (Montecino-Rodriguez *et al.* 1996). Thus, development of the immune system may be influenced by TR $\alpha$ , acting directly on B cells and via the cellular environment on T cells. Data from TR $\beta^{-/-}$  and TR $\alpha^{-/-}$  $\beta^{-/-}$  mice reveal decreased macrophage, granulocyte and T and B lymphocyte cellularity in TR $\beta^{-/-}$  mice, with a further reduction in TR $\alpha^{-/-}$  $\beta^{-/-}$  animals, indicating that lymphocyte development requires both TR $\alpha$  and  $\beta$  (Arpin *et al.* 2000).

Finally, Flamant *et al.* (2002) reported that TR $\alpha^{0/0}$  $\beta^{-/-}$  mice show a marked reduction in the size of the spleen, a finding that is more pronounced in Pax8 $^{-/-}$  mice. Pax8 $^{-/-}$  animals were observed to show a marked reduction in splenic follicle number and size. The spleen was also severely affected in Pax8 $^{-/-}$ TR $\beta^{-/-}$  double mutants (Flamant *et al.* 2002), whilst Pax8 $^{-/-}$ TR $\alpha^{0/0}$  double mutants showed a recovery of splenic phenotype (Flamant *et al.* 2002), suggesting that unliganded TR $\alpha$  isoforms, which may include the TR $\Delta\alpha$  variants, play an important role in spleen development.

## Summary

Analysis of the literature indicates that understanding the mechanism of T3 action *in vivo* has been advanced significantly by the generation of transgenic and knockout mice involving the TR $\alpha$  and  $\beta$  genes. These mice have provided detailed insight into the roles of the various TR isoforms. However, it is prudent to interpret data obtained from genetically modified mice with caution because of the unavoidable compensatory biological responses that occur following gene deletion or over-expression. Such responses may reveal phenotypes that are indirectly related to TR function or may mask important direct actions of TRs that can be performed by other, normally redundant, pathways. Furthermore, whilst comparisons between genetically modified mice can be very informative, an additional drawback is that different laboratories engineer mutations in mice with different genetic backgrounds.

Such differences may result in significant phenotypic modification. In future studies, therefore, it will be helpful to compare mutations of the TR $\alpha$  and  $\beta$  genes on uniform genetic backgrounds. In spite of these caveats, genetically modified mice have enabled T3-signalling pathways to be dissected in the intact animal. They have revealed the enormous complexity of TR function and provided a new and deeper understanding of the physiological actions of thyroid hormones.

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