

## Human Genome and Diseases: Review

### The TSH receptor and its role in thyroid disease

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**Abstract.** The thyrotropin (TSH) receptor plays a pre-eminent role in thyroid physiology and disease. TSH, acting through the TSH receptor, is the major stimulator of thyroid cell growth, differentiation and function. In Graves' disease, the TSH receptor is the target of stimulating antibodies that cause hyperthyroidism. Although still a topic of debate, the TSH receptor has been implicated in the pathogenesis of the endocrine ophthalmopathy associated with Graves' disease. Blocking antibodies against the TSH receptor are involved in the development of hypothyroidism in a subset of patients with autoimmune hypothyroidism. Transplacental passage of stimulating or blocking TSH receptor antibodies from a mother with autoimmune thyroid disease may result in transient hyper- or hypothyroidism in early infancy. During pregnancy, the placental hormone human chorionic gonadotropin

(hCG) can cause gestational hyperthyroidism through cross-reaction with the TSH receptor. Gestational hyperthyroidism may also be involved in the pathogenesis of hyperemesis gravidarum. Trophoblast tumors secreting hCG are a rare cause of hyperthyroidism. Somatic activating mutations of the TSH receptor have been identified as a molecular cause of toxic adenomas, whereas activating mutations in the germline give rise to nonautoimmune familial hyperthyroidism or sporadic congenital hyperthyroidism. These gain-of-function mutations are dominant, and one mutated allele is sufficient to result in disease. Inactivating germline mutations of both TSH receptor alleles lead to variable degrees of resistance to TSH, encompassing a spectrum ranging from euthyroid hyperthyrotropinemia to overt hypothyroidism with thyroid hypoplasia.

**Key words.** Thyroid; thyrotropin; receptor; cAMP; hyperthyroidism; hypothyroidism; autoimmune thyroid disease; mutation.

#### Introduction

The thyroid is controlled by a classic hypothalamic-pituitary axis (fig. 1). The hypothalamic tripeptide TRH (thyrotropin-releasing hormone) stimulates the production and secretion of the pituitary glycoprotein hormone TSH (thyroid-stimulating hormone). TSH is composed of an  $\alpha$  and a  $\beta$  subunit, the  $\alpha$  subunit being common to the glycoproteins TSH, FSH (follicle-stimulating hormone) and LH/CG (luteinizing hormone/chorionic gonadotropin). TSH, acting through its specific receptor, stimulates growth and function of thyroid follicular cells and regulates the synthesis and secretion of the thyroid hormones, thyroxine (T4) and triiodothyronine (T3) (fig. 1). Thy-

roid hormones have multiple effects on differentiation, growth and metabolism. Their action is primarily mediated by nuclear receptors regulating gene transcription, although nongenomic actions are increasingly recognized [1, 2].

In disease, the TSH receptor is directly involved in the pathogenesis of Graves' disease, autoimmune hypothyroidism, toxic adenomas, familial and sporadic nonautoimmune hyperthyroidism, and certain forms of resistance to TSH (fig. 2, 3). Not surprisingly, the TSH receptor has been a primary focus of interest, generating an impressive increase in the amount and complexity of knowledge since its cloning in 1989 [3]. The ample information on the many facets of the TSH receptor in physiology and

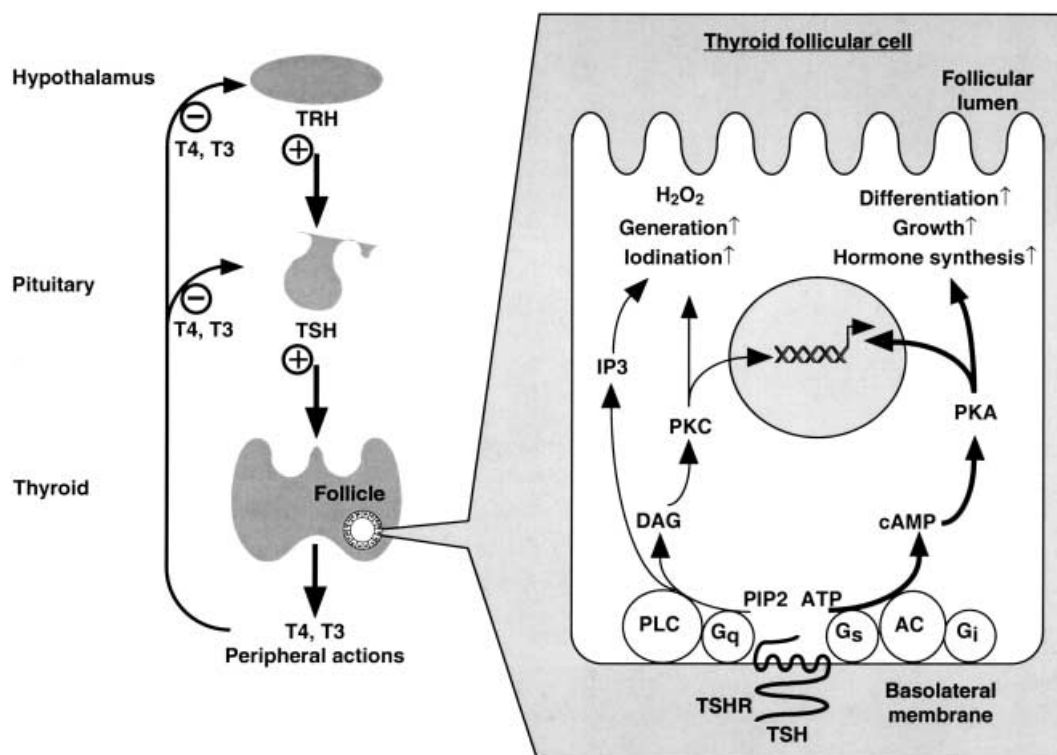


Figure 1. The hypothalamic-pituitary-thyroid axis and major signaling pathways in thyroid follicular cells. TRH, TSH releasing hormone; TSH, thyroid stimulating hormone; T4, tetraiodothyronine; T3, triiodothyronine; TSHR, TSH receptor; Gs/q/i, G proteins; AC, adenylate cyclase; PKA, protein kinase A; PLC, phospholipase C; DAG, diacylglycerol; PKC, protein kinase C; IP3, inositoltriphosphate.

disease is discussed in a number of reviews that contain further information and references not included here because of space constraints [4–18]. In addition to providing an overview of the implications of the TSH receptor in the pathogenesis of thyroid disorders, the primary emphasis of this review is directed towards the consequences of naturally occurring mutations in the TSH receptor.

### Characteristics of the TSH receptor

#### Gene and protein structure

The TSH receptor is a member of the large superfamily of G-protein-coupled seven-transmembrane receptors (GPCRs) (fig. 1, 4). It belongs to subfamily A, which is related to rhodopsin and the  $\beta_2$ -adrenergic receptor [19–21]. Together with the receptors for the other glycoprotein hormones FSH and LH/CG, the TSH receptor forms a distinct subgroup. The three glycoprotein hormone receptors share a high degree of homology (~70%) in the transmembrane domain [4, 6]. The main differences are found in the large amino-terminal extracellular domain involved in binding of the hormone. In the extracellular domain, the TSH receptor has 39% degree of homology with the FSH receptor, and 45% with the LH/CG receptor [4, 5, 8].

The canine TSH receptor complementary DNA (cDNA) was cloned using a reverse transcription polymerase chain reaction (RT-PCR) approach in 1989 [3, 4]. Mammalian TSH receptor cDNAs have subsequently been characterized in humans, rat, mouse, cow, sheep and cat [22–30]. Species comparisons reveal a high degree of homology of more than 90%. The single-copy human gene is located on chromosome 14q13, spans more than 60 kb and contains 10 exons [31–33]. The first 9 exons encode the majority of the extracellular domain. Exon 10 codes for the carboxy-terminal part of the extracellular domain, the whole transmembrane domain and the intracellular carboxy-terminus. The open reading frame consists of 2295 nucleotides encoding a 764-amino acid protein (fig. 4). A 21-amino acid signal sequence precedes the long extracellular domain of 397 amino acids. Splice variants of unknown physiologic relevance have been reported [34]. The TSH receptor differs from the LH/CG receptor by the presence of two unique insertions of 8 (residues 38–45) and 50 amino acids (residues 317–366) in the extracellular domain [4, 8, 14]. Deletion of residues 38–45 disrupts TSH binding, whereas deletion of residues 317–366 does not affect TSH binding or TSH-mediated cyclic AMP (cAMP) accumulation in transfected cells [35]. Chimeras of the TSH receptor with the LH/CG receptor reveal primary roles for the extracellular domain

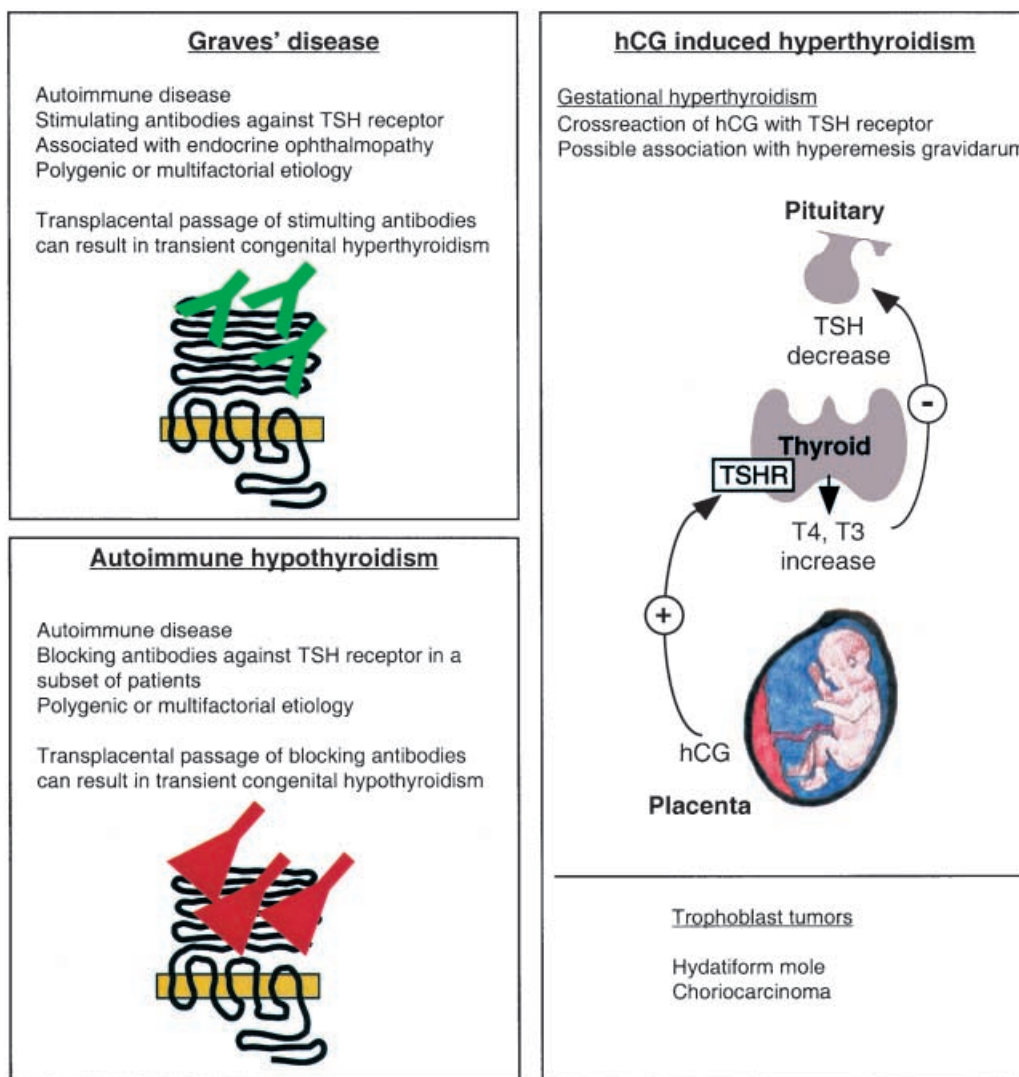


Figure 2. TSH receptor in disease without mutations.

in mediating the binding specificity, and for the transmembrane domain in signal transduction [36]. Currently, there is no information on the three-dimensional structure derived from crystallization of parts of the receptor, and expression of significant amounts of TSH-binding receptor remains a challenging task [14]. Based on the putative homology with the ribonuclease inhibitor, a three-dimensional model of the extracellular domain has been proposed [37]. Leucine-rich repeats are thought to form a horseshoe-shaped concave surface composed of parallel  $\beta$  sheets which interact with the ligand [7, 37, 38]. For the transmembrane domain, limited structural considerations have been proposed by comparing the TSH receptor to the helical wheel model proposed for GPCR and to a rhodopsin-derived model of the LH/CG receptor [39–42].

### Expression of the TSH receptor

In addition to its expression on the basolateral membrane of thyroid follicular cells (fig. 1), there is evidence for expression of the TSH receptor in several nonthyroidal tissues. These include fibroblasts, white and brown adipocytes, lymphocytes, certain regions of the brain, the adrenal glands, cardiac myocytes, the kidney and the thymus [29, 43–52]. Some of these observations are based on the detection of messenger RNA (mRNA) transcripts by RT-PCR, and this may have led to the detection of illegitimate transcripts [53]. However, convincing evidence for TSH receptor protein expression and function does exist for some of the tissues studied [14]. For example, neonatal human adipocytes display sensitivity to TSH [54], and retroorbital preadipocyte fibroblasts may differentiate into TSH receptor-bearing adipocytes under appropriate culture conditions [44, 55]. The TSH receptor

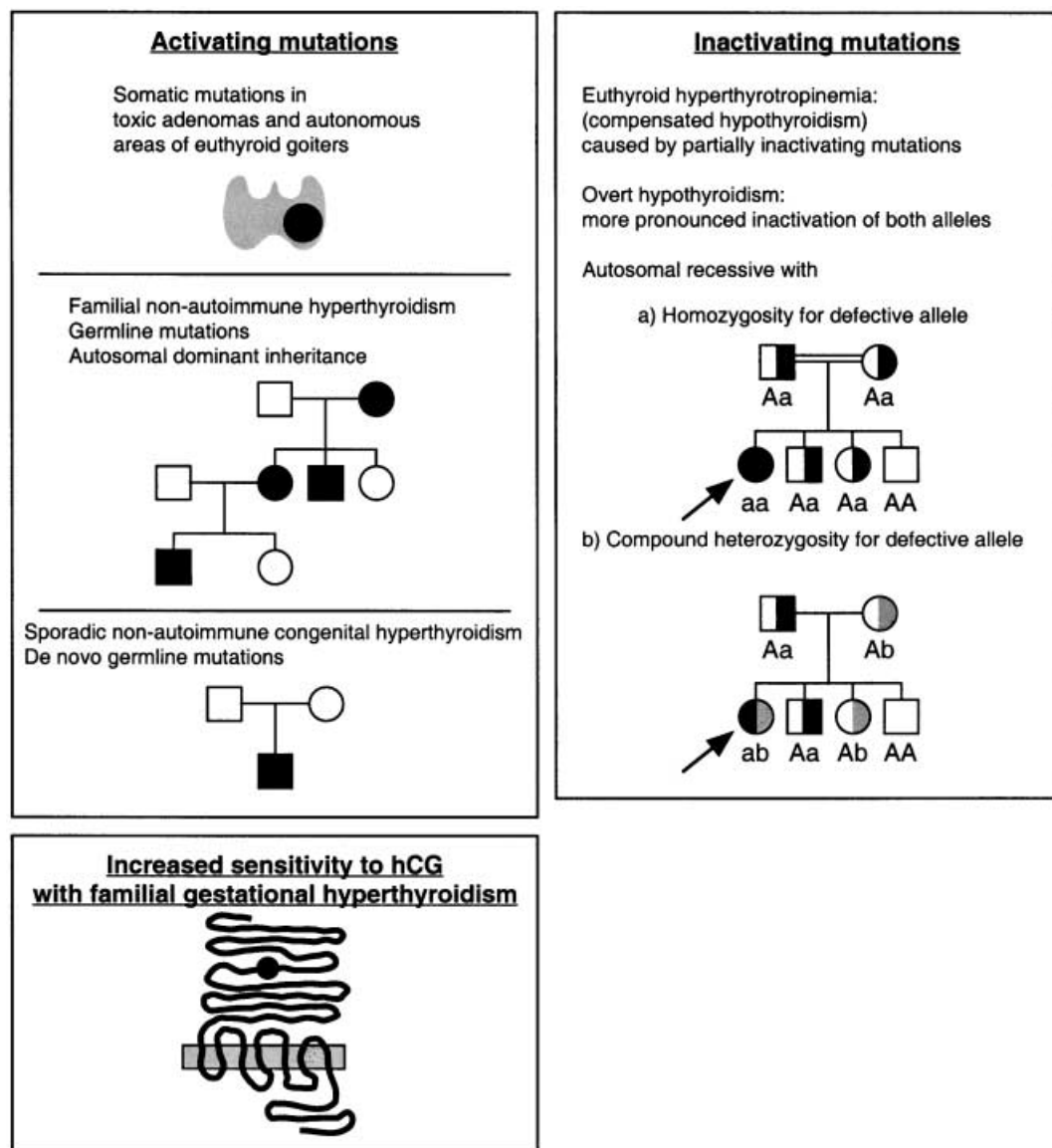


Figure 3. TSH receptor in disease with mutations.

cDNA has been cloned from ovine hypothalamus and appears to be expressed throughout the ovine brain [29]. In the anterior human pituitary, the TSH receptor was found to be expressed in folliculo-stellate cells, suggesting that it could be involved in an intrapituitary ultrashort loop regulating TSH secretion [48]. In the gastrointestinal tract, the TSH receptor has been detected in intestinal T cells but not in epithelial cells [46]. Interestingly, the *hyt/hyt* mouse, which is homozygous for an inactivating mutation in the TSH receptor, has a selectively impaired intestinal T cell repertoire, and it has been proposed that TSH may be a key immunoregulatory mediator in the intestine [46]. Dendritic cells were reported to have high levels of TSH receptors, and they display an enhanced phagocytic response as well as interleukin 1 and 12 secre-

tion after stimulation with TSH [47]. The functional relevance of TSH and its receptor in extrathyroidal tissues needs to be further delineated.

#### Glycosylation and palmitoylation

The human extracellular domain contains six potential asparagine-linked glycosylation sites (fig. 4). Based on studies in transfected CHO cells, the primary translation product of 85 kDa is glycosylated, resulting in a high-mannose form of ~100 kDa, with subsequent maturation to a 120-kDa form with complex carbohydrates [12, 14, 56]. TSH binding is increased on the mature, glycosylated form of the receptor, and mutation of glycosylation sites reduces TSH binding [12, 57]. The mature TSH re-

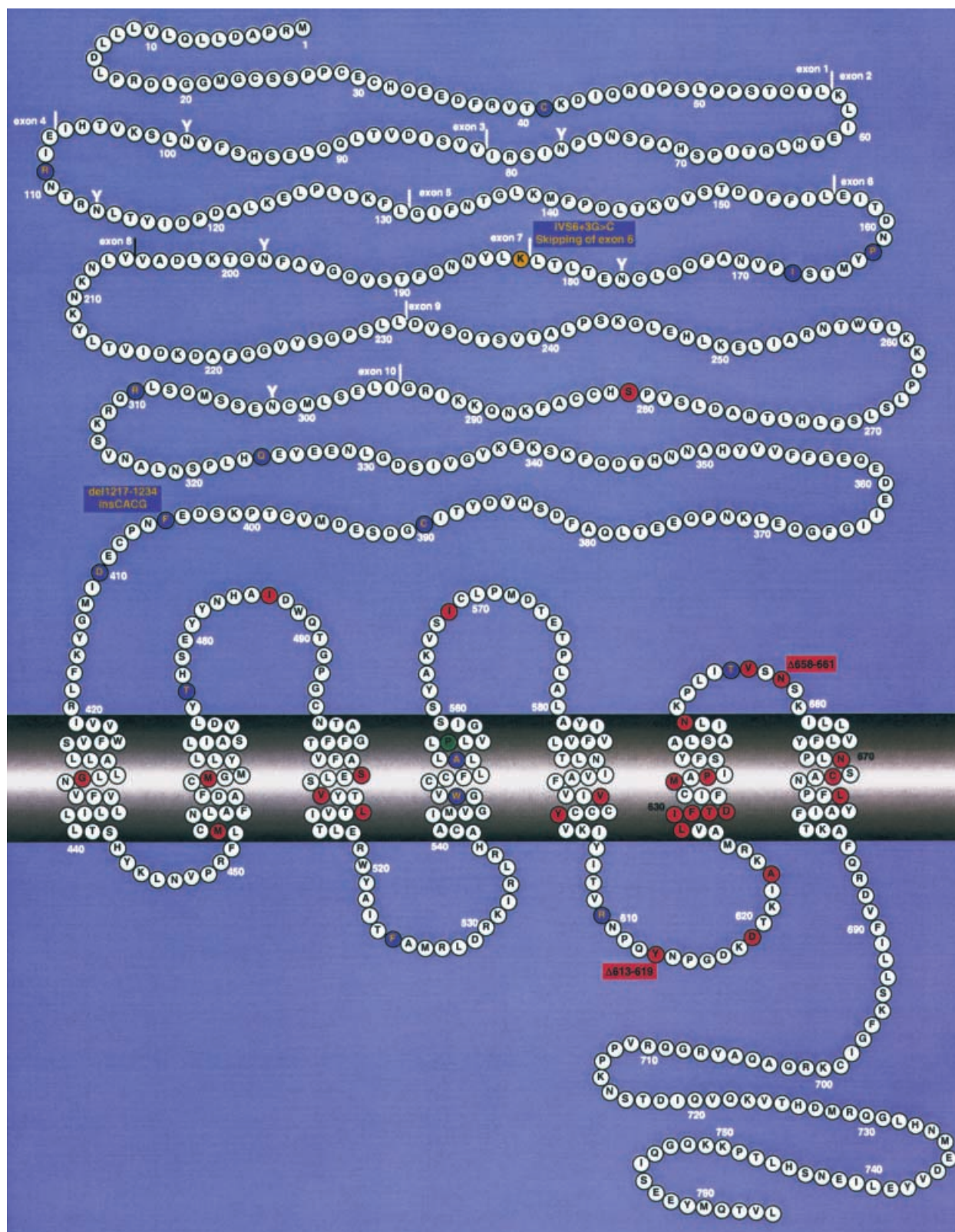


Figure 4. Naturally occurring mutations in the TSH receptor. ● Activating mutations, ● inactivating mutations, ● mutation conferring increased sensitivity to hCG, ● mutation in the hypothyroid *hyt/hyt* mouse. Y asparagine-linked glycosylation sites. The figure, and tables 1 and 2 include mutations published up to December 2000. The mutations are designated according to the *Recommendations for a nomenclature system for human gene mutations* [223].

ceptor is palmitoylated at cysteine 699 in the cytoplasmic tail of the receptor [58]. Mutation of cysteine 699 to alanine (C699A) delays surface expression, but the mutant remains fully active in terms of TSH binding and signaling.

### Two subunit model of the TSH receptor

Considerable interest has been directed towards the two-subunit model of the TSH receptor [14]. In vitro studies with thyroid cells in culture and mammalian cells transfected with the recombinant TSH receptor suggest that the receptor is present not only as a single chain, but also in a two-subunit form [59–61]. The glycosylated extracellular domain, the  $\alpha$  subunit, is cleaved from the unglycosylated membrane-spanning  $\beta$  subunit [14, 62]. The  $\alpha$  and  $\beta$  subunits (also referred to as A and B subunits) are linked by disulfide bonds [14]. The transmembranous  $\beta$  subunit is more prevalent in membrane preparations of thyroid follicular cells, suggesting that the  $\alpha$  subunit may undergo shedding from the transmembranous part of the receptor [61]. The cleavage sites that give rise to the two subunits flank the unique insertion found in the extracellular domain of the TSH receptor [17]. The amino-terminal cleavage site is located between residues 302 and 317 in proximity to a basic domain formed by residues 310–313 (RQR) [56, 63–65]. Deletion mutants of this region shift the cleavage site towards the amino terminus, suggesting that cleavage is not dependent on a specific amino acid sequence but rather occurs at a fixed distance from a protease attachment site [66]. The carboxy-terminal cleavage site seems to be located between residues 366 and 378 at or in proximity to a GQE motif at positions 367–369 [63–65, 67, 68]. The impossibility of recovering a connecting peptide suggests that the initial cleavage at the amino-terminal site is followed by progressive cleavage or degradation to the amino-terminal site or, alternatively, rapid disintegration of the putative connecting peptide [63–65, 67–69]. Initially, it was proposed that cleavage is performed by a metalloproteinase. More recent data, however suggest, that cleavage may be exerted by an enzyme that is related to TACE (tumor necrosis factor  $\alpha$ -converting enzyme) [64]. The enzyme is not thyroid specific since cleavage also occurs in transfected nonthyroidal cells expressing the TSH receptor.

The physiologic significance of the cleavage and shedding phenomenon remains unresolved. The existence of various isoforms generated from the initial holoreceptor raises the possibility that their biologic behavior is not uniform [56, 67]. However, affinity for TSH and signaling do not differ between the single-chain and the two-subunit forms of the receptor. Moreover, artificial TSH receptor mutations that prevent intramolecular cleavage do not abolish its constitutive activity [67]. Whether

shedding of the extracellular  $\alpha$  subunit or the release of fragments of the connecting peptide are involved in initiation of an immune response leading to the development of autoantibodies in Graves' disease remains unresolved [62].

### Desensitization

Analogous to many other GPCRs, the TSH receptor undergoes homologous desensitization after exposure to TSH [5]. Desensitization involves G protein receptor kinases (GRK 2 and GRK 5) which phosphorylate the receptor [70–72]. Subsequently,  $\beta$ -arrestin binds to the phosphorylated receptor, resulting in its uncoupling from  $G_{s\alpha}$  [71]. Transcriptional downregulation of the TSH receptor gene has been shown to involve a splice variant of CREM (cAMP response element modulator), the transcriptional repressor ICER (inducible cAMP early repressor) [73].

### Intracellular signaling in thyroid follicular cells

Binding of TSH to its receptor leads primarily to coupling to  $G_{s\alpha}$  and subsequent activation of adenylyl cyclase [74, 75] (fig. 1). At higher doses of TSH, the receptor couples to  $G_{q11}$ , resulting in activation of phospholipase C. Additional interactions have been demonstrated with other G protein families [76]. More recently, TSH-mediated phosphorylation of the intracellular Janus kinases (JAK) 1 and 2 and the JAK family substrate STAT3 (signal transducer and activators of transcription) has been demonstrated in rat thyroid cells and transfected Chinese hamster ovary (CHO) cells [77].

The TSH-dependent cAMP cascade is the major regulator of growth, differentiation and hormone secretion of thyroid follicular cells [78]. The increase in cAMP leads to phosphorylation of protein kinase A and to activation of targets in the cytosol and the nucleus, such as the transcription factor CREB. Stimulation of  $G_{q11}$  and the phospholipase C-dependent inositol phosphate/diacylglycerol pathway at higher doses of TSH activates hydrogen peroxide generation and iodination [74, 75, 79, 80]. In contrast to the receptors for LH/CG and FSH, the TSH receptor displays constitutive basal activity, i.e. readily measurable spontaneous activity in the absence of ligand. The predominant role of the cAMP pathway in thyroid cell growth and function has been corroborated by several transgenic models with chronic overstimulation or disruption of this pathway [81–84]. Its importance is furthermore underscored by the consequences of antibodies that stimulate or block the TSH receptor, as well as naturally occurring activating and inactivating mutations of the TSH receptor [13].

## The TSH receptor in disease

Disorders affecting thyroid function or growth are common and may occur independently or in combination. Functional disturbances result in hyper- or hypothyroidism. In iodine-sufficient regions, the autoimmune thyroid disorders, Hashimoto thyroiditis (hypothyroidism) and Graves' disease (hyperthyroidism), are by far the most common clinical entities affecting the thyroid gland. Their prevalence is significantly higher in women [85, 86]. Benign growth alterations are equally common and include diffuse and multinodular goiter, cysts, nodules and adenomas. Thyroid carcinomas are relatively infrequent neoplasms that account for ~0.6–1.6 % of all malignancies [87]. Figures 2 and 3 summarize the disorders in which the TSH receptor plays a pivotal role in the pathogenesis.

### Graves' disease

Graves' disease is a complex autoimmune disease. Its exact pathogenesis, which involves T and B cells, remains unresolved, and a discussion of these aspects is beyond the scope of this review. In addition to the signs and symptoms of thyrotoxicosis, patients with Graves' disease often develop Graves' ophthalmopathy, and they may present with skin infiltrations, most commonly pretibial myxedema.

In Graves' disease, stimulatory antibodies (TSABs) directed against the TSH receptor, the major antigen in this form of autoimmune thyroid disease, mimic TSH action and result in hyperthyroidism and goiter [88, 89] (fig. 2). Antibodies against the TSH receptor are heterogeneous. In addition to TSAB, blocking antibodies (TBAB) may be present concomitantly, and fluctuations in their relative concentrations may result in periods of hyper- or hypothyroidism [90, 91]. Routine assays do not differentiate between TSAB and TBAB, but merely measure the binding of immunoglobulins to the receptor by a radioreceptor assay using porcine or human TSH receptor (TBI, TSH-binding inhibition or TBIs, TSH-binding inhibitory immunoglobulins). More sensitive second-generation TBI assays use a monoclonal-antibody-capturing recombinant TSH receptor in a coated tube format [92]. Determination of TBIs can be useful in the differential diagnosis of hyperthyroidism, and they can be helpful in the prediction of fetal and neonatal thyroid dysfunction in neonates born to a mother with a history of autoimmune thyroid disease. There is also some suggestion that TBI levels may be useful as predictors in the course of Graves' disease [91]. Functional characterization of TRAb is possible using thyroid cells or cell lines stably transfected with the human TSH receptor [91]. Most commonly these assays rely on the determination of intracellular cAMP accumulation. More recent developments include

measurement of cAMP-dependent luciferase reporter gene activity [93, 94].

The TSH receptor epitopes for TSAB and TBAB are not well defined [9, 14, 17]. Studies with linear epitopes suggest that TSABs bind predominantly to the amino-terminal part of the extracellular domain, whereas TBABs bind primarily to an ectodomain region in proximity to the transmembrane domain [8, 95]. However, it is more likely that the majority of these autoantibodies recognize conformational epitopes that involve segments of both the amino- and carboxy-terminal segments of the ectodomain [14, 96]. Chimeras of the TSH and LH/CG receptor not only confirm that autoantibodies interact with the TSH receptor ectodomain but provide evidence that TSH and autoantibody-binding sites are overlapping, but not identical [96].

Although several studies suggested that certain naturally occurring sequence variations of the TSH receptor may have immunogenic properties or result in enhanced signaling activity, it seems unlikely that any of these polymorphisms is of relevance in the pathogenesis of Graves' disease (for review see [9]). The possibility that autoantibodies against the TSH receptor arise in response to a foreign antigen with structural similarity, for example infection with *Yersinia enterocolitica*, has been raised, but remains unresolved [14, 97].

A genetic component in the pathogenesis of Graves' disease is supported by family and twin studies [98, 99]. Given the key role of TSABs in the pathogenesis of Graves' disease, the TSH receptor gene itself is an obvious candidate gene. Most population-based case-control studies argue, however, against a significant role of the TSH receptor locus [100–102], although a positive association has been reported in one study on women with Graves' disease [103]. Linkage studies and transmission equilibrium analyses have equally been negative [102, 104]. Susceptibility loci that may be involved in the pathogenesis of Graves' disease currently include the *HLA* locus (6p21), the *CTLA-4* gene (2q33), and loci referred to as *GD-1* (14q31), *GD-2* (20q11.2) and *GD-3* (Xq21.33–22) [99].

### Ophthalmopathy and skin manifestations in Graves' disease

Graves' disease is often associated with ophthalmologic symptoms, endocrine ophthalmopathy or Graves' ophthalmopathy, and skin manifestations [105–107]. It is important to differentiate signs of increased sympathomimetic activity associated with all forms of thyrotoxicosis from the tissue involvement defining endocrine ophthalmopathy. The latter includes edema, chemosis, congestion of conjunctival vessels, proptosis, extraocular muscle involvement (diplopia), corneal lesions due to lagophthalmos and compression of the optic nerve [106]. Minor ocular involvement is present in the majority of

patients more pronounced ophthalmopathy occurs in ~25–50% [106]. Skin manifestations are most commonly seen in the pretibial area, but Graves' dermopathy may occur in any area of the body in response to trauma or pressure [107].

Because of the fact that the TSH receptor is the major antigen in Graves' disease, and given the well-established role of TSABs in the pathogenesis of hyperthyroidism and goiter, these molecules form obvious candidates in the pathogenesis of eye and skin disease. Reports on the identification of TSH receptor mRNA transcripts in orbital tissues from patients with endocrine ophthalmopathy by RT-PCR were met with some skepticism due to concern that these messages may be illegitimate transcripts without physiologic relevance [53, 108, 109]. However, TSH receptor mRNA has also been detected using a ribonuclease protection assay in orbital tissue from patients with Graves' disease, but not from normal controls, suggesting that the receptor is indeed expressed in this tissue [44]. Moreover, retroorbital preadipocyte fibroblasts appear to differentiate into TSH receptor-bearing adipocytes under appropriate culture conditions [55]. These studies suggest that the TSH receptor may indeed be expressed in orbital tissue in Graves' disease. However, it remains unclear whether the TSH receptor is the orbital antigen responsible for the development of endocrine ophthalmopathy. There are currently no studies that document reactivity of orbital T cells against the TSH receptor, and, as of yet, the autoantigen(s) and specific pathogenic autoantibodies have not been identified.

The documented expression of the TSH receptor in adipocytes and fibroblasts may also be of relevance for the development of skin manifestations [14, 110, 111]. Interaction of autoantibodies with the TSH receptor or a cross-reacting antigen would, however, not explain the predilection of extrathyroidal manifestations in retroorbital tissue and pretibial skin, nor the occurrences at other locations in response to trauma. Chronic inflammation, in combination with local physical factors, may therefore be required for the development of the extrathyroidal manifestations of Graves' disease [111].

#### **Animal models for Graves' disease**

Graves' disease probably only occurs in humans. Cats are the only other mammalian species in which spontaneous thyrotoxicosis occurs frequently [112]. The pathogenic mechanisms underlying feline hyperthyroidism remain controversial, as both an autonomous mechanism of growth and function, and an autoimmune etiology have been proposed [112, 113]. However, a recent study, using cells transfected with the feline TSH receptor, provides further evidence against the presence of TSH receptor antibodies in hyperthyroid cats [30].

Initial attempts to create murine models of Graves' disease by immunization with soluble TSH receptor and ad-

juvant were not successful in eliciting appropriate T cell and B cell responses, and histological analyses only revealed lymphocytic infiltrates in a subset of these experiments [14]. However, several promising approaches in generating mouse models of Graves' disease have been generated during the last few years [16, 114–120]. For example, the immunization of mice with fibroblasts stably transfected with cDNA encoding the human TSH receptor and major histocompatibility complex (MHC) class II antigen resulted in thyrotoxicosis and goiter in ~25% of the animals [115]. The sera contained detectable stimulating antibodies against the TSH receptor, as determined in CHO cells expressing the TSH receptor. However, there was no lymphocytic infiltration in thyroid tissue and no ocular involvement [115]. Inbred mice immunized with a plasmid encoding the human TSH receptor developed TBII and thyroiditis in the majority of the animals [117]. This approach was also successful in inducing lymphocytic and mast cell infiltration of the orbits [118]. Moreover, there was accumulation of adipocytes and edema in the orbital tissue, periodic acid Schiff-positive material, probably corresponding to glycosaminoglycans, dissociation of muscle fibers and TSH receptor immunoreactivity. This interesting model may be a first example of endocrine ophthalmopathy caused by the transfer of T cells sensitized to the TSH receptor, and it supports the concept that this disease has indeed an autoimmune basis [118]. Particularly promising is a recent model with outbred mice immunized with human TSH receptor cDNA in a eukaryotic expression vector [120]. All mice developed antibodies capable of recognizing the recombinant TSH receptor in CHO cells. Consistent with the higher prevalence of Graves' disease in women, 17 of 29 females, but only 1 of 30 males, developed overt hyperthyroidism, goiter with lymphocytic infiltration and ocular signs reminiscent of Graves' ophthalmopathy [120].

#### **Graves' disease during pregnancy**

Hyperthyroidism is second to diabetes mellitus as the most common endocrinopathy in pregnancy. Its prevalence is ~0.05–0.2% [121, 122]. Its clinical assessment may be difficult, because many of the symptoms suggestive for hyperthyroidism are also associated with normal pregnancy. Hyperthyroidism during pregnancy is most commonly caused by Graves' disease or hCG-induced gestational hyperthyroidism (discussed below). The natural history of Graves' disease in pregnancy is typically characterized by aggravation during the first trimester, amelioration in the second and third trimesters, and recurrence a few months after pregnancy [123]. In mothers with autoimmune thyroid disease, the fetus may rarely present with congenital hyperthyroidism caused by maternal-to-fetal transfer of thyroid-stimulating antibodies. This form of hyperthyroidism is transient because the



stimulating antibodies are cleared from the fetal circulation. Very rarely, congenital hyperthyroidism is caused by activating mutations in the TSH receptor, and in these infants it is persistent (see below) [124].

### **Human chorionic gonadotropin-induced gestational hyperthyroidism and hyperemesis gravidarum**

During early pregnancy, secretion of the placental hormone human chorionic gonadotropin (hCG) may result in thyroid dysfunction. At high doses, hCG cross-reacts with the TSH receptor, and this stimulation can lead to an increase in secretion of T4 and T3, with subsequent suppression of TSH secretion [123, 125] (fig. 2). The increased secretion of hCG may result exclusively in suppression of TSH, or in overt hyperthyroidism. Estimates for hCG-induced hyperthyroidism predict that it occurs in ~30–40 patients in 1000 pregnancies [123]. Elevations of hCG are particularly pronounced in twin pregnancies [126]. Because of the decrease in the levels and bioactivity of hCG later in pregnancy, this form of hyperthyroidism is typically transient and limited to the first 3–4 months of gestation. Gestational hyperthyroidism is frequently associated with hyperemesis gravidarum [123, 127]. The hyperemesis might be caused by a marked hCG-induced increase in estradiol levels [128]. However, the relation between hyperemesis and gestational hyperthyroidism varies among patients, and additional, unidentified mechanisms may be involved. Hyperthyroidism is only rarely caused by trophoblastic tumors, hydatiform moles and choriocarcinomas that secrete high amounts of hCG [129]. In men, choriocarcinomas can arise in the testis and cause hyperthyroidism by secreting hCG [130].

### **Hypersensitivity to hCG due to mutation of the TSH receptor**

A remarkable form of familial gestational hyperthyroidism caused by a mutant thyrotropin receptor displaying hypersensitivity to normal levels of hCG has been identified by Rodien et al. (table 1; fig. 3, 4) [131]. The proband had a history of two miscarriages that were accompanied by hyperemesis. Subsequently, she had two pregnancies that were complicated by hyperthyroidism, severe nausea and vomiting. Of note, she was negative for antibodies against the TSH receptor or thyroid peroxidase (TPO). Her hCG levels, determined during the second pregnancy, were in the normal range for the first trimester. The patient's mother had a history of one miscarriage and two pregnancies that were complicated by hyperemesis gravidarum. Direct analysis of the TSH receptor gene in the proband and her mother revealed the presence of a heterozygous point mutation in exon 7, resulting in the substitution of K183R (fig. 4). Functional studies in COS-7 cells transfected with the mutated receptor documented no dif-

ferences in membrane expression, and similar levels of basal and TSH stimulated cAMP accumulation. Whereas the wild-type TSH receptor reacts only minimally to high doses of hCG, the K183R mutant is hypersensitive to hCG, although it is still 1000 times less responsive to hCG than the LH/CG receptor. Aside from explaining the recurrent hyperthyroidism in these two patients, the K183R TSH receptor mutation is unique because sensitivity is increased for hCG but remains unaltered for the cognate ligand TSH [131]. This observation also supports the possibility of a hCG-independent connection between hyperthyroidism and hyperemesis gravidarum.

### **Autoimmune hypothyroidism**

In adults living in iodine-replete regions, autoimmune thyroiditis is the most common cause of hypothyroidism [85, 86]. Analogous to Graves' disease, it is a complex autoimmune disorder with an abnormal humoral and cellular immune response to the thyroid. Several forms of autoimmune thyroiditis are distinguished on clinical grounds [132]. In Hashimoto's thyroiditis, the thyroid shows the signs of widespread chronic lymphocytic infiltration, and at least in early stages of the disease, it is mildly enlarged [133]. In addition to an increase in TSH and a decrease in thyroid hormone levels, most patients have elevated titers of autoantibodies against thyroperoxidase (Anti-TPOAb) and thyroglobulin (Anti-TGAb) [133]. The progressive destruction of thyroid tissue by chronic inflammation may ultimately lead to a decrease in thyroid hormone synthesis and hypothyroidism. In about 15% of patients with autoimmune hypothyroidism, the thyroid dysfunction is thought to be the consequence of blocking antibodies directed against the TSH receptor (fig. 2) [134–136]. Of note, transplacental passage of TBAb may be a cause of transient congenital hypothyroidism [137]. Linkage and association studies indicate a role for the HLA and CTLA-4 genes in the development of autoimmune hypothyroidism, but the TSH receptor locus itself has not been implied as a susceptibility locus [138].

### **Activating mutations in the TSH receptor**

The discovery that site-directed mutagenesis of a critical residue in the third intracellular loop of the  $\alpha_{1b}$ -adrenergic receptor can lead to constitutive activation in the absence of ligand subsequently led to the search and detection of naturally occurring activating mutations in numerous GPCRs [139, 140]. Gain of function mutations are by definition dominant, and alteration of one allele is thus sufficient for generating the pathologic phenotype. Activating TSH receptor mutations can occur somatically in solitary toxic adenomas or toxic adenomas within multinodular goiters. Germline TSH receptor mutations give rise to autosomal dominant nonautoimmune hyperthy-

Table 1. Activating TSH receptor mutations.

Residue	Toxic adenoma <sup>a</sup>	Thyroid cancer <sup>b</sup>	Familial nonautoimmune hyperthyroidism <sup>b</sup>	Sporadic congenital hyperthyroidism <sup>b</sup>
<i>Activating somatic and germline mutations</i>				
S281	S281I S281T S281N			S281N [194, 186]
G431			G431S [186]	
M453	M453T	M453T [181]		M453T [179, 180]
M463			M463V [187]	
I486	I486F I486M	I486F [175]		
S505			S505R [39] V509A [155]	S505N [193, 196]
V509				
L512	L512R			
I568	I568T			
V597				V597L [195]
Y601	Y601N			
613-619	Del613-619 YNPGDKDTK			
D619	D619G			
A623	A623I A623S A623V	A623V [171]	A623V [183]	
L629	L629F		L629F [184]	
I630	I630L			
F631	F631L F631C			F631L [124]
T632	T632I T632A	T632I [172] T632A [172]		T632I [192]
D633	D633A T633Q T633Y D633H	D633H [173]		
M637	M637R			
P639	P639S		P639S [185]	
N650			N650Y [39]	
V656	V656F			
658-661	Del658-661 NSK			
N670			N670S [39]	
C672			C672Y [155]	
L677		L677V [174]		
<i>Increased sensitivity to hCG</i>				
Residue	Familial gestational hyperthyroidism <sup>b</sup>			
K183	K183R [131]			

<sup>a</sup> The reports on TSH receptor mutations in toxic adenomas are too numerous to cite individually. The interested reader is referred to the following articles and reviews [7, 10, 13, 146].

<sup>b</sup> References are indicated in brackets.

roidism or, in the case of de novo mutations, to sporadic nonautoimmune congenital hyperthyroidism (table 1; fig. 3, 4).

### TSH receptor mutations in toxic adenomas

Chronic stimulation of the cAMP cascade results in enhanced proliferation and function of thyroid follicular cells [78]. Consequently, any molecular alteration leading to constitutive activation of the cAMP pathway in a thyroid cell is predicted to result in clonal autonomous

growth and function, and ultimately in a toxic adenoma [78]. Such somatic mutations were discovered in the stimulatory  $G_{s\alpha}$  subunit (*gsp* mutations) in toxic adenomas, and have also been found in a subset of nonfunctioning adenomas and differentiated thyroid carcinomas [141–143].  $G_{s\alpha}$  mutations, most commonly found in amino acids arginine 201 and glutamine 227, impair the hydrolysis of GTP to GDP, resulting in persistent activation of adenylyl cyclase. The same  $G_{s\alpha}$  mutations are found in 35–40% of somatotroph tumors in acromegalic

patients [144]. Mosaicism for  $G_{\alpha s}$  mutations that occur early in development cause the McCune Albright syndrome and may affect multiple tissues, including the thyroid [145].

Somatic mutations in the TSH receptor were first discovered in toxic adenomas [146]. The first identified mutations were clustered in the third intracellular loop and the sixth transmembrane domain of the receptor, but a wide variety of activating somatic mutations have been found in subsequent studies (table 1; fig. 4) [7, 10, 13, 147]. Mutations conferring constitutive activity occur in the entire transmembrane domain, including the extracellular loops, as well as the carboxy-terminal region of the extracellular domain. All these mutations increase basal cAMP levels, but only a few amino acid substitutions activate the phospholipase C cascade in a constitutive manner [13]. Inositoltriphosphate (IP<sub>3</sub>) accumulation in response to TSH is, however, usually retained. A naturally occurring mutation in the fifth transmembrane domain, Y601N, is distinct in this respect in that it results in constitutive activation of the cAMP pathway, while losing coupling to  $G_{\alpha 11}$  [148]. A polymorphic variant at this position, Y601H, loses basal cAMP accumulation while retaining responsiveness to TSH, and it also fails to activate the PLC pathway [41, 148].

The reported prevalence of TSH receptor mutations in toxic adenomas varies widely, but may be as high as 80% [147, 149]. Differences in sampling technique and methodological approach, as well as variations in iodine intake, may contribute to the reported differences in prevalence rates [150, 151].

The vast majority of activating mutations is located in the  $\beta$  subunit of the receptor. However, naturally occurring substitutions at serine 281 (S281I/N/T), a residue located in the extracellular  $\alpha$  subunit, also confer constitutive activation to the receptor (fig. 4) [152, 153]. The exact mechanisms resulting in gain of function are still poorly defined. Some of the mutations may alter the positions of the transmembrane helices, thus mimicking the conformational changes induced by binding of ligand. Alternatively, some mutations may alter the structure of domains that inhibit receptor coupling to G proteins in the absence of TSH [154, 155]. Activating mutations in the extracellular domain are thought to result in the relief of a negative constraint present in the unliganded carboxy-terminal part of the extracellular domain [147, 152, 153, 156, 157], and a subset of mutations may alter desensitization of the receptor.

### TSH receptor mutations in multinodular goiters

Multinodular goiters are common, particularly under conditions of scarce iodine supply, and there is an age-dependent increase in their prevalence [85]. The mechanisms underlying nodular alteration are complex [158],

and the following discussion is limited to hyperfunctioning adenomas within multinodular goiters or autonomous areas within euthyroid goiters. Somatic TSH receptor mutations have been identified in toxic adenomas isolated from multinodular goiters [159–161]. Consistent with studies demonstrating distinct clonal origins of different thyroid adenomas within the same multinodular goiter [162], the mutations may differ among these adenomas. For example, in two adenomas from the same goiter, one neoplasm harbored a mutation M453T, the second adenoma a T632I substitution [159]. In another study, L632I and F631L were found in two distinct lesions within the same goiter, whereas another patient had two distinct toxic nodules with the same mutation (I630L) [160]. These studies demonstrate that the pathogenesis of hyperfunctioning adenomas does not differ between multinodular goiters and solitary toxic adenomas. In contrast, TSH receptor mutations are rare in nonfunctioning adenomas, even if of monoclonal origin [161, 163]. Therefore, distinct mechanisms must be implicated in the abnormal growth leading to nonfunctioning nodules [158].

A germline polymorphism in the cytoplasmic tail of the receptor, D727E, has been associated with the development of multinodular goiters [164]. This polymorphism was found in 8 of 24 patients, and *in vitro* studies by these authors suggested an increased cAMP accumulation of the D727E variant in response to TSH. These findings contrast with a study characterizing basal and TSH-stimulated cAMP-dependent transcriptional activity using a luciferase reporter gene that was unable to document a difference between the wild type and the D727E variant [165]. In a more recent study, the association between the codon 727 polymorphism and toxic thyroid adenomas or toxic multinodular goiters could not be confirmed [166]. In a study analyzing hyperfunctioning and nonfunctioning areas from patients with toxic multinodular goiters, activating TSH receptor mutations were detected in 14 of 20 hyperfunctioning areas, whereas no mutation was identified in nonfunctioning nodules [167]. On microscopic analysis, only two of the hyperfunctioning areas corresponded to classic adenomas surrounded by a capsule, whereas the remainder had the characteristic features of hyperplastic lesions. The clonal composition of the analyzed tissues was not determined, and it has to be emphasized that histologic analysis does not allow to discriminate monoclonal from polyclonal lesions [168]. Constitutively activating TSH receptor mutations have also been detected in autoradiographically hyperfunctioning areas of goiters from euthyroid patients [169]. These findings contribute to the understanding of the mechanisms underlying thyroid autonomy and functional heterogeneity in multinodular goiters [167, 169].

### TSH receptor mutations in thyroid carcinomas

In well-differentiated thyroid cancers, mutations in the  $G_{sa}$  subunit and the TSH receptor genes occur only rarely [143, 170–175]. Although constitutive activation of the cAMP pathway results in enhanced growth, it is not thought to be sufficient for malignant transformation of otherwise normal thyrocytes.

Interesting aspects have been gained from the study of a patient presenting with hyperthyroidism and increased uptake in two nodules, but suppressed uptake in the remainder of the gland [173]. After surgical removal of the right thyroid lobe, histological examination revealed the presence of a papillary carcinoma of the insular subtype with lymph node metastases. On follow-up examination with a whole-body scan, this patient had diffuse uptake in both lungs, but no uptake in residual tissue in the neck. Therefore, the diagnosis of autonomously functioning thyroid cancer was made. Mutational analysis of the TSH receptor gene documented a somatic mutation, D633H, located in the sixth transmembrane domain, in DNA isolated from the primary tumor and metastatic tissue. This aspartate residue is conserved among the glycoprotein hormone receptors. In the LH receptor, the corresponding mutation D578H has been found in Leydig cell tumors [176]. The D578H LH receptor displays constitutive activation of both the cAMP and phospholipase C (PLC) signaling pathways in transfected cells [176]. Recently, the substitution of this aspartate to histidine has been analyzed in more detail in the LH, TSH and FSH (D581H) receptors [177]. Similar to the D578H LH receptor, the D633H TSH receptor produces strong constitutive activation of both the cAMP and PLC pathways. In the FSH receptor, D581H does not spontaneously stimulate inositol phosphate production, but it confers the ability to mediate FSH-stimulated PLC activity, a property absent in the wild-type FSH receptor. PLC stimulation is pertussis toxin insensitive, indicating that a member of the Gq family mediates the response. No other amino acid that has been substituted for this conserved aspartate in helix 6 of glycoprotein hormone receptors has the same functional impact as histidine. This suggests that this histidine stabilizes a receptor conformation with increased ability to couple to Gq. The ability of the D578H LHR and D633H TSH receptor mutants to costimulate the cAMP and PLC pathways constitutively may thus have direct relevance to the pathogenesis of endocrine neoplasia [177]. Another mutation that activates both pathways, I486F, was found in a hyperfunctioning well-differentiated follicular carcinoma in a patient presenting with hyperthyroidism and increased radioiodine uptake within the thyroid mass (table 1). This supports the notion that concomitant activation of these two signaling cascades may promote transformation [175]. In a patient with a Hürthle cell carcinoma, Russo et al. identified a L677V TSH receptor mutation [174]. Basal cAMP levels were increased in

transfected CHO cells, but IP3 accumulation has not been determined with this mutant.

Overexpression of the naturally occurring M453T mutation in the FRTL-5 rat cell line was sufficient to induce neoplastic transformation as assessed by growth in semi-solid medium and athymic mice [178]. This mutation only activates the cAMP pathway in transfected COS (green monkey fibroblast-like) cells, and the neoplastic transformation in FRTL-5 cells may be linked to inherent properties of this cell line, which is known to be tumorigenic under conditions of prolonged TSH stimulation. In humans, this mutation has been found in the germline of two patients with congenital hyperthyroidism, but there was no suggestion that it is oncogenic on its own (table 1) [179, 180]. However, a somatic M453T substitution has been identified in a 11-year-old girl with a hyperfunctioning nodule and a papillary carcinoma [181]. This raises the possibility that this mutation could itself be associated with neoplasia, or that one or several mutations in other growth-controlling genes have occurred in cells harboring the M453T mutation.

### Germline mutations in familial nonautoimmune hyperthyroidism

Autosomal dominant familial hyperthyroidism without evidence of an autoimmune etiology was first recognized by Thomas et al. [182]. The typical signs associated with autoimmune hyperthyroidism, i.e. exophthalmos, myxedema, TSAbs and lymphocytic infiltration of the thyroid gland, are absent in this form of hyperthyroidism. Because all thyroid follicular cells display an increased growth rate, these patients have a diffuse goiter. The molecular basis of inherited nonautoimmune hyperthyroidism was elucidated by detecting activating germline mutations in the TSH receptor in the family reported by Thomas et al. [155], and subsequently confirmed in several other families (table 1; fig. 3, 4) [39, 183–187]. The onset of hyperthyroidism may vary in carriers of the same mutation in a given kindred. Thus, other factors, for example genetic background and/or iodine intake, appear to modulate the phenotypic expression [182, 188, 189]. In a Chinese family with a P639S germline mutation and hyperthyroidism in the father and three children, two of the children and the father also had mitral valve prolapse (MVP) associated with mitral regurgitation [185]. The close temporal relationship between the onset of thyrotoxicosis and the diagnosis of mitral valvular disease was taken as an indication that an increased clinical expression of MVP may exist in genetically predisposed individuals.

### Sporadic germline mutations in congenital hyperthyroidism

Congenital hyperthyroidism is usually caused by transplacental passage of stimulating TSH receptor autoantibodies [190]. Autoimmune neonatal hyperthyroidism is

rare and occurs in less than 2% of infants born to a mother with a history of Graves' disease [191], a condition with an estimated incidence of about 2 of every 1000 pregnancies [182]. Antibody-induced neonatal hyperthyroidism typically resolves within the first 3–7 months as the maternal antibodies are cleared from the circulation. A fluctuating course may be seen because of the concomitant presence of TBABs [90].

Constitutively activating *de novo* mutations of the TSH receptor have been found in a few patients with sporadic congenital nonautoimmune hyperthyroidism (table 1; fig. 3, 4) [124, 179, 180, 186, 192–196]. Congenital hyperthyroidism due to a toxic adenoma harboring a somatic TSH receptor mutation was reported as another unusual variant of congenital hyperthyroidism [153]. These rare cases with nonautoimmune congenital hyperthyroidism due to TSH receptor mutations must be differentiated from the much more frequent and transient autoimmune form of hyperthyroidism, because most of these patients have pronounced hyperthyroidism requiring a more aggressive therapeutic approach that may necessitate surgery and ablative radiotherapy. It is noteworthy that several of these children with severe neonatal hyperthyroidism seem to have mild mental retardation [124, 192, 193], suggesting that high levels of thyroid hormone may have a negative impact on brain development [197]. A subset of these children had proptosis [179, 180]. Computertomography of the retroorbital tissue in one of these children failed, however, to demonstrate infiltration of the eye muscles [179].

#### **Inactivating mutations in the TSH receptor**

Resistance to TSH may be caused by various molecular mechanisms. In pseudohypoparathyroidism type 1a (PHP 1a), the inability of TSH to stimulate the cAMP pathway is the consequence of inactivating mutations in  $G_{s\alpha}$ . PHP1a is typically associated with resistance to other hormones such as gonadotropins. Moreover, these patients present with characteristic somatic features that include short stature, brachydactyly and round face (Albright's hereditary osteodystrophy) [198]. Resistance to TSH can also occur as an isolated phenomenon without the features of PHP 1a (for review see [11, 15]). In a subset of these patients, the molecular cause consists of inactivating mutations in the TSH receptor that are partially or completely inactivating (table 2; fig. 3) [199–206]. The mode of inheritance is recessive and affected individuals are homozygous or compound heterozygous for inactivating mutations. Among these patients the phenotype encompasses a wide spectrum ranging from isolated TSH elevation to severe hypothyroidism, and there is a clear correlation between genotype and phenotype. In other patients with sporadic or familial resistance to TSH, the TSH receptor gene was found to be normal, indicating

locus heterogeneity due to defects in other genes [207–209]. Obvious candidate genes include genes encoding elements of the TSH-dependent signaling cascades or regulators of thyroid development and gene expression.

#### **The *hyt/hyt* mouse**

A loss-of-function mutation in the TSH receptor gene as a cause of TSH resistance was first discovered in the hypothyroid *hyt/hyt* mouse [26, 27]. The phenotype of this inbred mouse strain is defined by congenital hypothyroidism, retarded growth, mild anemia, hearing loss and infertility [210–214]. The mode of inheritance is autosomal recessive. The hypoplastic thyroids of these mutant mice are located in the proper position. Histologically, the thyroid follicular cells are developed, but incompletely differentiated, and the epithelial cells are not organized into structures recognizable as follicles [211]. The thyroid tissue of homozygous mice is unresponsive to TSH *in vivo* and *in vitro*, but activators of  $G_{s\alpha}$  induce a normal cAMP response [214]. Analysis of the mouse TSH receptor cDNA ultimately led to the identification of a missense mutation resulting in the substitution of a highly conserved proline (P556L) in the fourth transmembrane domain by leucine (fig. 4) [26, 27]. The mutation eliminates TSH binding, and thus normal receptor function, although the membrane localization of the receptor appears to be preserved [27].

#### **Euthyroid hyperthyrotropinemia caused by TSH receptor mutations**

The first human case with TSH resistance due to a defect in the TSH receptor was documented in three sisters, offspring of unrelated parents, who were found to have normal peripheral thyroid hormone but high TSH levels, a constellation referred to as euthyroid hyperthyrotropinemia (table 2) [199]. Both parents only showed discrete TSH elevations. None of the family members had clinical signs of hypothyroidism. The three affected siblings were found to be compound heterozygous for mutations in the extracellular TSH binding domain of the receptor (P162A and I167N) (table 2; fig. 4) [11, 199]. *In vitro* studies documented that the I167N mutation had almost no biologic activity, whereas P162A displayed reduced activity [199]. Cell surface expression of the P162A mutant was reduced about twofold, and the EC50 for TSH stimulation was increased twofold [38]. In contrast, the I167N mutant did not reach the cell surface. Both amino acids are part of a putative  $\alpha$  helix formed by the fifth leucine rich repeat motif in the extracellular domain and are thought to influence interaction of the receptor with the ligand [7, 11, 37]. Based on the current model of the three-dimensional structure of the TSH receptor, the P162A substitution maps to the surface of the molecule, whereas the I167N mutation affects a residue whose side chain contributes to

Table 2. Inactivating TSH receptor mutations.

Protein	Gene <sup>a</sup>	Membrane expression	Basal and TSH-induced cAMP	Thyroid size	Thyroglobulin	Comments	Ref.
<i>Euthyroid hyperthyrotropinemia</i>							
P162A	484C > G	reduced	shifted to the right	normal	not reported	three affected sisters	38, 199
I167N	500T > A	retained intracellularly	almost none				
C41S	122G > C	studied in [224]	studied in [224]	normal <sup>123</sup> I-iodine scan	normal		201
F525L	155C > A	expressed on membrane, TSH binding similar to wt	reduced basal and TSH stimulated activity				
Q324X	970C > T	not analyzed	not analyzed	slightly enlarged on <sup>123</sup> I-iodine scan	normal		201
D410N	1228G > A	expressed on membrane, TSH binding similar to wt	reduced basal and TSH stimulated activity				
C390W	1170T > G	expressed on membrane, TSH binding reduced	basal activity similar to wt reduced response to TSG	slightly enlarged on <sup>123</sup> I-iodine scan	normal		201
W546X	1637G > A	not analyzed	not analyzed				
P162A	484C > G	studied in [38, 199]	Studied in [38, 199]	Normal on <sup>123</sup> I-iodine scan	normal	consanguinity	201
R310C	928C > T	TSH binding reduced	constitutive activity no response to TSH	normal ultrasound and scan	not reported	consanguinity discrepancy in vitro and in vivo (see text)	215
<i>Hypothyroidism</i>							
R109Q	326G > A	TSH-binding 60% of wt	right-shifted response to TSH	normal ultrasound and scan	not reported	mild hypothyroidism	200
W546X	1637G > A	no specific binding	inactive				
Frameshift at F406 > 420X	Del1217-1234/ins CACG	retained intracellularly inserted into membrane	absent	hypoplastic on ultrasound	normal off levothyroxine	mild hypothyroidism	202
C390W	1170T > G	total binding similar to wt	reduced basal levels, right-shifted response to TSH				
A553T	1657G > A	Normal translation, very low levels of membrane expression $K_d$ reduced	reduced basal and TSH-induced cAMP accumulation no coupling to Gq11	hypoplastic on ultrasound Tc-scan negative	high normal range	consanguinity profound hypothyroidism	203
Altered splicing with skipping of exon 6 T656X	IVS6 + 3 G > C	not reported	not reported	hypoplastic on ultrasound TC-scan negative	normal range	profound hypothyroidism maternal great aunt with congenital hypothyroidism of distinct etiology	204
R609X	1825C > T	not reported	not reported	hypoplastic on ultrasound normal location Tc-scan negative	below or within the normal range	profound hypothyroidism hypocortisolism consanguinity ACTH resistance, reason unknown	205
T4771	1430C > T	very low cell surface expression receptor retained in intracellular compartments $K_d$ reduced	no basal activity no significant increase in response to TSH	hypoplastic on ultrasound <sup>131</sup> I-iodine scintigraphy: reduced volume, low uniform uptake	undetectable	profound hypothyroidism common ancestors	206

<sup>a</sup> The mutations are designated according to the *Recommendations for a nomenclature system for human gene mutations* [223].

the hydrophobic core characteristic of proteins harboring leucine repeat motifs [37, 38]. I167N seems to cause a gross destabilization of receptor structure that is incompatible with its normal routing through the intracellular membrane system of the cell [38].

Euthyroid hyperthyrotropinemia due to mutations in the TSH receptor gene has subsequently been reported in several other families (table 2; fig. 4) [201, 215]. The size of the thyroid gland is either normal or slightly increased in these patients. Partial resistance to TSH associated with a homozygous R310C replacement was found in two brothers with compensated hypothyroidism [215]. In transfected cells, this mutation has reduced TSH binding and absent responsiveness to TSH, but it appears to have constitutive activity in terms of cAMP accumulation in comparison to the wild-type receptor. If constitutive activity of the R310C mutations is indeed responsible for the compensated hypothyroid state, it is not clear why TSH levels were elevated in the two patients homozygous for the mutation.

#### **Congenital hypothyroidism due to TSH receptor mutations**

More pronounced or complete inactivation of both TSH receptor alleles leads to mild or severe congenital hypothyroidism (table 2; fig. 3, 4) [202–206]. Because of absent uptake of the radioisotope, scintigraphic studies typically do not reveal any thyroid tissue. However, ultrasound of the neck will reveal the presence of a normally located hypoplastic gland. Intriguingly, many of these patients have normal or elevated thyroglobulin levels. For example, in an inbred kindred studied by Abramowicz et al., the hypothyroid index patient had no detectable thyroid tissue on scintigraphy prompting the diagnosis of thyroid agenesis [203]. His thyroglobulin levels were, however, in the high normal range, and ultrasonography documented a normally located hypoplastic thyroid gland. The patient and his sister, who had been diagnosed earlier with hypothyroidism, were found to be homozygous for a A553T mutation in the fourth transmembrane domain (table 2; fig. 4). Functional analysis of the mutated receptor in transfected cells demonstrated an extremely low level of expression of the mutated receptor at the cell surface despite normal intracellular synthesis. The mutated receptor was able to bind TSH and stimulate cAMP accumulation in transfected cells, but appeared to have lost the ability to activate the PLC pathway. The finding of high thyroglobulin levels in this situation is not entirely clear. It suggests that thyroglobulin measurements, combined with ultrasound of the neck, may be helpful in distinguishing congenital athyreosis from thyroid hypoplasia, an entity which may eventually escape detection by scintigraphy [203, 204].

In other patients with inactivation of both TSH receptor alleles, thyroglobulin levels were undetectable or low

(table 2). For example, in a female patient with severe hypothyroidism, thyroid hypoplasia and a homozygous mutation in the first extracellular loop (T477I) with loss of function, thyroglobulin levels were undetectable [206]. In a large inbred Bedouin kindred in which profound congenital hypothyroidism and hypocortisolism occurred alone or together in eight family members, serum thyroglobulin levels varied among the affected individuals and were below or within the normal range [205]. The patients were homozygous for an inactivating mutation truncating the TSH receptor in the third intracellular loop (R609X). The molecular cause of the adrenal insufficiency remained unclear, but was not linked to the ACTH receptor. The presence of hypothyroidism did not affect the timing, severity and manner of clinical manifestation of hypocortisolism [205].

#### **Significance of inactivating TSH receptor mutations for thyroid development**

The *hyt/hyt* mouse and patients with TSH-resistant congenital hypothyroidism with correctly located hypoplastic glands confirm that development and migration of the thyroid is independent of TSH stimulation [202, 203, 211]. This is consistent with the observation that the genes for thyroperoxidase, thyroglobulin and the TSH receptor are only expressed once the gland has reached its pretracheal location [216, 217]. Although early events of thyroid development are not dependent on TSH and its signaling pathway, this cascade is essential for complete differentiation, growth and function of thyroid follicular cells [26, 84, 211].

Whereas congenital hypothyroidism is a relatively frequent disorder affecting ~1:3000 newborns, thyroid hypoplasia is only found in about 5% of all patients with congenital hypothyroidism and inactivating mutations in the TSH receptor only accounts for a subset of these patients. A study of familial or consanguineous cases of congenital hypothyroidism was negative for linkage to the TSH receptor locus [218]. Moreover, mutational analysis of the coding sequence of the TSH receptor did not reveal sequence alterations that would explain the phenotype in several studies of patients presenting with thyroid hypoplasia, hypothyroidism and resistance to TSH [201, 207–209]. It has also become apparent that hypoplasia can be associated with several genetic defects, including mutations in the TSH $\beta$  subunit, the TSH receptor and the G $_{sa}$  subunit. More recently, mutations in the paired domain transcription factor PAX8 have been identified as a further cause of thyroid hypoplasia [219]. This is supported by the murine knockout model; mice homozygous for a disrupted *Pax8* gene have a phenotype with hypothyroidism, severely hypoplastic thyroid glands and absence of follicular structures [220]. Homozygosity for mutations in the forkhead transcription factor TTF2 (thyroid transcription factor 2; FKHL15) has been demonstrated

in two siblings with thyroid agenesis, cleft palate, choanal atresia and spiky hair [221]. Similarly, targeted disruption of *Ttf2* in the mouse leads to cleft palate and either a sublingual or completely absent thyroid gland [222].

## Perspective

Although an overwhelming wealth of new insights into the function of the TSH receptor in physiology and disease has been obtained, numerous challenging aspects remain to be solved or refined [4, 6, 14]. They include, among others, further clarification of the exact role of the TSH receptor in the pathogenesis of Graves' disease and endocrine ophthalmopathy, further refinement of the current animal models for Graves' disease, development of new assay systems for the determination of autoantibodies directed against this receptor and last but not least, attempts to elucidate parts of its structure.

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