

The Biology of pro-Thyrotropin-Releasing Hormone-Derived Peptides*

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I. Introduction

A SIGNIFICANT body of research has led to the conclusion that peptides are important regulatory components of nervous system function. Initially, the “peptidergic neuron” concept was reserved for the neurosecretory cells in the hypothalamus that released oxytocin and vasopressin directly into the circulation from their nerve termi-

nals in the posterior pituitary. The idea of neurosecretion in the hypothalamus can be traced back to the work of Scharrer and Scharrer (1) as early as the late 1920s. Later work by Harris (2) specified that the hypothalamic substances secreted into the portal vessels were pituitary specific and led to the concept of “releasing factors” whose purpose was to initiate a cascade of events resulting in the release of peripherally active hormones (3). The discovery and chemical characterization of the first identified hypothalamic releasing factor, TRH (pyro-Glu-His-ProNH₂, also known as thyroliberin, and herein referred to as TRH) by Guillemin and colleagues (4) and Schally and colleagues (5) provided ultimate confirmation for the founding principles of neuroendocrinology and was followed by the discovery of other peptide-releasing factors (6, 7).

TRH, produced in the paraventricular nucleus of the hypothalamus (PVN),¹ stimulates the biosynthesis and secretion of TSH from the anterior pituitary (8, 9). TSH, in turn,

¹ Abbreviations: iTRH, immunoreactive TRH; TRH-OH, deamidated TRH; CHP, cyclo (His-Pro); PTU, propylthiouracil; HPT, hypothalamic-pituitary-thyroid; DVC, dorsal vagal complex; DMN, dorsal motor nucleus of the vagus; NTS, nucleus tractus solitarius; IML, intermediolateral column of the spinal cord; SP, substance P; NPY, neuropeptide Y; CGRP, calcitonin gene-related peptide; CT, calcitonin; IL1, interleukin 1; VIP, vasoactive intestinal peptide; OAG, the diacylglycerol analog 1-oleoyl-2-acetyl-sn-glycerol; CCK, cholecystokinin; NT, neurotensin; THC, tetrahydrocannabinol; ACh, acetylcholine; ECS, electroconvulsive shock treatment; NE, norepinephrine; E, epinephrine; DA, dopamine; 5-HT, serotonin; NO, nitric oxide; NMDA, N-methyl-D-aspartate; 6-HODA, 6-hydroxydopamine; 5,7-DHT, 5,7-dihydroxytryptamine; GABA, γ -aminobutyric acid; EOPs, endogenous opioid peptides; CA, catecholamines; ENK, enkephalins; DYN, dynorphin; SRIF, somatostatin; CP, carboxypeptidases; CPE, carboxypeptidase E; CSF, cerebrospinal fluid; NAc, nucleus accumbens; VTA, ventral tegmental area; PVN, paraventricular nucleus of the hypothalamus; RPa, nucleus raphe pallidus; RMg, nucleus raphe magnus; ROB, nucleus raphe obscurus; icv, intracerebroventricular; sc, subcutaneous; ic, intracerebral; SHR, spontaneously hypertensive rats; WKY, Wistar-Kyoto rats; MCA, middle cerebral artery; ALS, amyotrophic lateral sclerosis; RMN, Rolling mouse Nagoya model of ataxia; CNS, central nervous system; EEG, electroencephalogram; PAG, periaqueductal gray; RPi, nucleus reticularis paraventricularis; MAF, mesencephalic reticular formation; POA, preoptic nucleus of the hypothalamus; LC, locus coeruleus; LPGi, nucleus paraventricularis lateralis; LS, long-sleep; SS, short-sleep; PBMC, peripheral blood monocyte; ME, median eminence; RER, rough endoplasmic reticulum; PCs, proconverting enzymes; RSP, regulated secretory pathway; GC, Golgi complex; SG, secretory granules; TGN, trans-Golgi network; ICC immunocytochemistry; TRE, thyroid response element; CRE, cAMP response element.

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stimulates thyroid hormone biosynthesis and release (10). TRH is central in regulating the hypothalamic-pituitary-thyroid (HPT) axis. TRH influences the release of other hormones, including PRL, GH, vasopressin, and insulin (11–13), and the classic neurotransmitters noradrenaline and adrenaline (14). Further, TRH is present in many brain loci outside of the hypothalamus, supporting a potential role as a neuromodulator or neurotransmitter outside of traditional HPT axis function (15, 16). For example, TRH is implicated as a modulator of seizure activity (17) and gastrointestinal function (18). TRH has been also found outside the central nervous system (CNS) in the gastrointestinal tract, pancreas, reproductive tissues including placenta, ovary, testis, seminal vesicles, and prostate, retina, and blood elements (19). The widespread distribution of TRH within and outside the CNS supports a diverse range of roles for this molecule, roles likely to involve many functions outside of the traditional HPT axis.

Of the many peptide products derived from the TRH precursor (pro-TRH described below), until recently only TRH itself was studied extensively. In the last few years, a new wave of research has identified many other products derived from pro-TRH and suggests potential biological functions for these non-TRH peptides. The immediate precursor to TRH, TRH-Gly, independent of conversion to TRH, stimulates gastric acid secretion in a dose-dependent manner, although it is 100-fold less potent than TRH (18). Other peptides derived from pro-TRH, originally referred to as cryptic peptides because their roles are incompletely understood, are now being studied. In the hypothalamus, prepro-TRH_{160–169} (pST₁₀, also known as Ps4) and prepro-TRH_{178–199} (pFE₂₂), peptides that lie between the third and fourth and the fourth and fifth progenitor sequences for TRH in the TRH precursor, respectively, are released from perfused rat hypothalamic slices and the median eminence (20). [The reader should note that in Fig. 1, pro-TRH-derived peptides are named by “p” for peptide followed by the single letter amino acid designation for the first and last amino acid of the peptide, along with the peptide length in subscript. Where these peptides are first mentioned, they are followed by the longer prepro-TRH name that describes their amino acid residue positions within the precursor.] prepro-TRH_{160–169} potentiates TSH release from anterior pituitary and stimulates TSH β gene promoter activity (21). This peptide also potentiates TRH-induced gastric acid secretion when microinjected into the dorsal motor nucleus of the vagus (22). Thus, prepro-TRH_{160–169} acts in concert with TRH both within and outside traditional HPT roles. prepro-TRH_{178–199} is proposed to be a CRH-inhibiting factor (Refs. 23 and 24 but see Ref. 25).

Both prepro-TRH_{25–50} (pYE₂₆) and prepro-TRH_{53–74} (pFT₂₂) are released from the median eminence (ME) and isolated anterior pituitary cells in response to depolarizing concentrations of potassium (26). In general, pro-TRH-derived peptides are present in the ME in much higher levels than other regions of the brain, consistent with complete processing of pro-TRH in the ME (E. A. Nillni, unpublished data). *In vivo* studies demonstrate that pro-TRH processing is regulated (27). During opiate withdrawal in rats, prepro-TRH mRNA is induced in the periaqueductal gray (PAG). The N-terminal peptides prepro-TRH_{53–74} and propro-

TRH_{83–106} (pEH₂₄) are increased (see Section IV), whereas the level of TRH remains unaltered (27, 28). In addition, prepro-TRH_{178–199} and its processed forms are increased in the PVN during suckling (see Section IV). Thus, levels of various products derived from pro-TRH can be independently regulated under altered physiological conditions.

Substantial progress has been made in the last few years in understanding the biosynthesis and processing of prohormone and neuropeptide precursors. For pro-TRH (Fig. 1), we have developed a model of its processing to mature peptides (29–31). Like many other secreted peptides, processing of the primary translation product, prepro-TRH, begins with removal of the signal peptide during its passage into the lumen of the rough endoplasmic reticulum (RER). From our current knowledge, processing of pro-TRH takes place within the regulated secretory pathway (RSP) (for full definition see Section IIA). Two recently discovered serine proteases, which are members of the family of prohormone convertases (PCs), PC1 (SPC3) and PC2 (SPC2), related to subtilisin and the yeast-processing enzyme Kex 2 (32–34), are the primary PCs involved in posttranslational processing of pro-TRH (31, 35–37).

Recent contributions to the understanding of pro-TRH biosynthesis and processing provide a useful framework for uncovering the diversity of function displayed by pro-TRH-derived peptides and the way that this diversity is generated. We will review the current knowledge of pro-TRH processing to the most studied biological end product, TRH, and importantly, other pro-TRH-derived peptides. The importance of understanding pro-TRH processing is then underscored by a comprehensive review of the wide range of potential biological roles subsumed by non-TRH products derived from pro-TRH. Although data are limited, where possible we summarize where these peptides are produced and how their levels might be regulated. We then review the HPT axis, with emphasis of its effects on pro-TRH processing. This is followed by a comprehensive review of the function of extrahypophysiotropic TRH. A review of TRH and other pro-TRH-derived peptide receptors then explains mechanisms of pro-TRH peptide signal transduction. Finally, TRH degradation provides an additional way by which peptide levels are tightly controlled, and this area is reviewed. The degradation of other pro-TRH-derived peptides is not yet understood.

II. Biosynthesis of TRH and Other pro-TRH-Derived Peptides

A. Biosynthesis and processing of pro-TRH

In recent years considerable research has focused on the expression of neuropeptide genes and their tissue-specific regulation. However, it has become clear that the peptides derived from these genes play significant neuromodulatory roles in the control of the CNS neurotransmitters. Even more astounding is the discovery that multiple neuropeptides with distinct physiological functions arise from the processing of single polypeptide precursors (36, 38–40). Thus, to fully understand the biology of a neuropeptide, one must understand the processing of the prohormone gene prod-

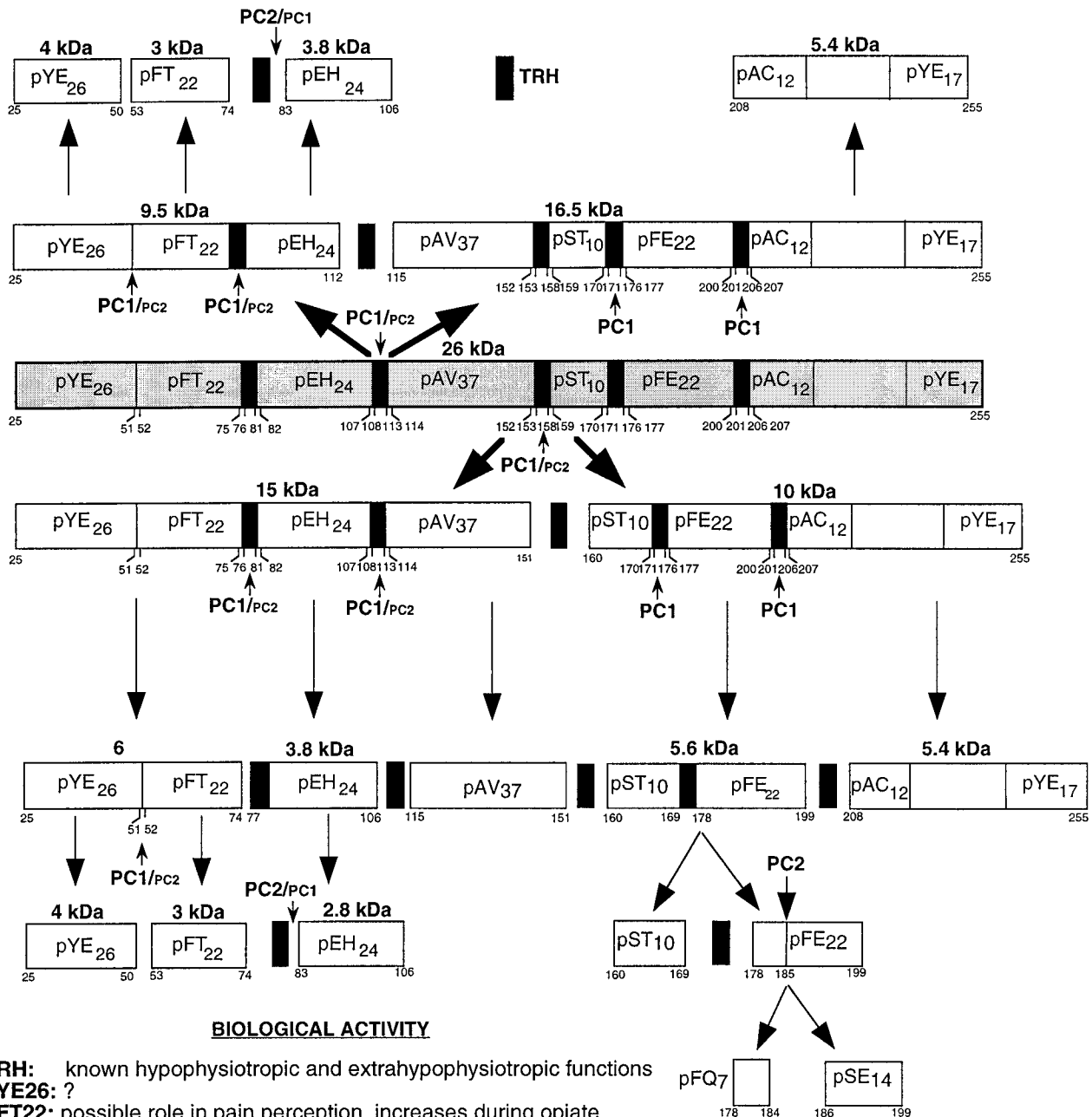


FIG. 1. Diagrammatic representation of the proposed processing model of rat pro-TRH to non-TRH peptides and TRH, and the postulated involvement of PC1 and PC2 as determined from our studies (for further details see Section II.A-D). The *small arrows* indicate PC1 and PC2 activity. Cleavage sites and direction of the processing cascade are indicated with *longer arrows*. *Thicker arrows* indicate that most of the initial cleavage of the intact precursor was produced at this site. The positions of paired basic residues are indicated by numbers. Non-TRH peptides are indicated in the *shaded* pro-TRH molecule, and TRH is indicated by a *black rectangle*.

uct, as well as the regulation of the gene's transcription. How do cells produce different levels of one peptide with respect to another when both derive from the same protein se-

quence? This is achieved through differential processing and degradation by the action of specific enzymes acting in specific cellular and extracellular compartments. Posttransla-

tional processing of hormone precursor proteins is a critical mechanism by which cells increase their biological and functional diversity, such that two or more peptides with different biological functions originate from the same precursor. It is through differential posttranslational processing mechanisms that cells selectively produce specific peptides for secretion.

Even though the amino acid structures of vasopressin and insulin had been elucidated in the 1950s (41), it was not until the early 1960s that the mechanisms of protein biosynthesis began to be understood and the genetic code was fully defined. In 1964, Sachs and Takabatake (42) provided the first evidence that the biosynthesis of vasopressin can be inhibited with puromycin, a protein synthesis inhibitor, and that newly synthesized vasopressin could not be detected in tissues until more than 1 h after pulse labeling. In these studies, Sachs and Takabatake demonstrated that before vasopressin becomes a biologically active peptide, it exists in a modified, or proform, state. Posttranslational modification was then required to convert the proform into an active peptide.

While the structure of insulin peptides was described early on (41), it was difficult to envision how the combination of A and B chains was attained in β -cells of the pancreas. Experiments initiated by Steiner in 1965 using tritiated leucine and phenylalanine to label proinsulin and insulin represented a landmark in prohormone theory. Using a pancreatic insulinoma derived from a patient, it was possible to determine that insulin could be derived *in vitro* from a single molecule that was converted to the A and B chains by trypsin treatment (43). Studies done in rat islets subsequently demonstrated conversion of proinsulin to insulin, in a relatively slow process taking approximately 40 min (44). During the same period, work done by Howell and Taylor (45) on insulin biosynthesis showed that newly synthesized insulin was released several hours after its biosynthesis. The emerging view of these findings was that some sort of orderly vectorial transport occurred involving the RER, the Golgi complex (GC), and secretory granules (SG). These data (46), along with the major contributions of other investigators who established that the biosynthesis of serum albumin, PTH, and glucagon also originate from larger precursors, formed the basis of the prohormone theory. This theory states that synthesis of peptide hormones and neuropeptides begins with mRNA translation into a large, inactive precursor peptide, followed by limited posttranslational proteolysis to release bioactive end products. Chretien and Li (47) also made an important contribution to the prohormone theory when they determined the amino acid sequences of β -lipotropin (β -LPH), γ -LPH, and β -melanotropin (β -MSH). They observed that β -MSH was part of the β -LPH sequence, providing evidence that β -MSH was a conversion product of β -LPH. They also observed that cleavage occurred at the C-terminal side of paired basic lysine or arginine residues. More definitive evidence for a precursor/product was provided with the cloning of POMC (29 kDa), which revealed that the ACTH and β -LPH sequences were present within the N terminus of POMC (38).

In summary, the biochemical processing of neuropeptides, peptide hormones, and other secreted proteins begins with limited posttranslational proteolysis of larger inactive pre-

cursors. Prohormones generally have their hormone sequences flanked by a single, a pair or four (tetra) basic amino acids where subtilisin-like processing enzymes produce their initial endoproteolytic cleavage (48, 49). This endoproteolytic cleavage is produced at the C-terminal side of the single or paired basic amino acid residue(s) which is followed by removal of the basic residue(s) by carboxypeptidase enzymes (CP) (50, 51). Further modifications can occur in the form of N-terminal acetylation, pyroglutamate formation, or C-terminal amidation, which confers bioactivity to many peptides (52).

At the intracellular level, hormone precursors are synthesized on membrane-bound ribosomes, by which they are translocated into the lumen of the RER via a signal recognition peptide. During vectorial transport through the GC and beyond, the newly synthesized proteins are subjected to posttranslational modifications including glycosylation, phosphorylation, amidation, acetylation, and proteolytic conversion (48). Ultimately, partially processed proteins reach the last compartment of the GC, the *trans*-Golgi Network (TGN). At the TGN, unprocessed or partially processed products are sorted to the RSP (29, 53, 54) to be stored in immature secretory granules (ISG). Upon maturation, electron-dense SGs containing sorted products can fuse with the plasma membrane in response to an extracellular stimulation in a calcium-dependent manner, thereby releasing their contents into the external milieu (55). Two pathways of unstimulated release are proposed for AtT₂₀ cells: constitutive (nongranular) secretion and basal release from compartments that form after sorting into the RSP (56). The mechanism whereby constitutive and regulated proteins are differentially sorted into separate vesicles after budding from the TGN is still under intensive investigation (57, 58).

Two hypotheses have been proposed to explain how proteins are selectively targeted from the TGN to the RSP. In the first hypothesis, proteins are sorted by passive aggregation, in which the proteins condense within forming ISG, thereby excluding other proteins from entering in the granule. This process occurs under acidic pH and high calcium concentrations (59, 60). Support for the aggregation hypothesis comes from studies done with chromogranin A (61, 62), chromogranin B (62), carboxypeptidase E (CPE) (63), and prohormone convertase 2 (PC2) (64). However, there are data suggesting that aggregation alone is not sufficient for sorting. Modifications of the chromogranin B sequence (65) can prevent the correct sorting of these peptides to the RSP, while their *in vitro* aggregation properties appeared unaltered. The insulin-like growth factor-1 (66) does not aggregate in the TGN, but is still sorted in the RSP.

The second hypothesis, originally proposed in 1985 by Kelly (67), involves *cis*-acting sorting signals within a protein destined for the RSP that interact with membrane-bound sorting receptors. Sorting receptors, possibly located in the forming SG, direct segregation of the protein for further packaging into SGs. Protein aggregation within the ISG can occur in this model, but is more critical for product concentration than sorting *per se*. Evidence supporting this second hypothesis has come from experiments involving chimeric proteins (68–73), where the fusion of constitutively secreted protein to a protein destined for RSP caused a rerouting of

this protein to the RSP. Conversely, proteins that have their sorting signal domains modified may be misrouted from the RSP into the constitutive pathway, as demonstrated for POMC (74), chromogranin A and B (75), PC2 (69), and glycine α -amidating monooxygenase (PAM) (76). A related hypothesis is sorting by retention, in which all proteins are initially targeted to ISGs, after which proteins that do not belong in the RSP are removed to their final destination, *e.g.*, lysosomal enzymes (57, 77).

It has been proposed recently that the membrane form of CPE, localized to the TGN, is a sorting signal receptor (78). CPE is proposed to direct POMC, proinsulin, proenkephalin, but not chromogranin A, into the RSP (79). Thus, CPE is a common sorting receptor for some, but not all, prohormones, and there must be other sorting receptors to direct trafficking of other proteins to the RSP (79). However, Irminger *et al.* (80) have provided evidence refuting the claim that CPE is a sorting receptor for proinsulin. In those studies they used pancreatic islets isolated from CPE-deficient (*Cpe^{fat}/Cpe^{fat}*) and control (*Cpe^{fat}/+*) mice to examine whether the trafficking of proinsulin and insulin was affected. They found that CPE was not essential for the sorting of proinsulin to the RSP (80). However, similar experiments with procholecystokinin in *Cpe^{fat}/Cpe^{fat}* mice indicate that CPE does function as a sorting receptor (81).

With this as a background, we turn specifically to the processing of pro-TRH. The elucidation of the rat prepro-TRH sequence in 1986 was of key importance to understanding the processing of pro-TRH to TRH and non-TRH peptides. Rat prepro-TRH is a 29-kDa polypeptide composed of 255 amino acids. The rat precursor contains an N-terminal 25-amino acid leader sequence, 5 copies of the TRH progenitor sequence Gln-His-Pro-Gly flanked by paired basic amino acids (Lys-Arg or Arg-Arg), 4 non-TRH peptides lying between the TRH progenitors, an N-terminal flanking peptide, and a C-terminal flanking peptide (82, 83). The N-terminal flanking peptide (prepro-TRH₂₅₋₅₀-R-R-prepro-TRH₅₃₋₇₄) is further cleaved at the C-terminal side of the arginine pair site to render prepro-TRH₂₅₋₅₀ and prepro-TRH₅₃₋₇₄, thus yielding a total of 7 pro-TRH-derived peptides (Fig. 1).

The biosynthesis of TRH and other pro-TRH-derived peptides follows the same prohormone-processing mechanisms described above, beginning with mRNA-directed ribosomal translation, followed by posttranslational limited proteolysis of the larger precursor, proTRH. This process occurs while pro-TRH is transported from the TGN to newly formed ISGs (84). These granules then mature and are targeted to sites of secretion at the plasma membrane of the cell. Rat, mouse, and human pro-TRH, similar to other peptide hormone precursors such as pro-enkephalin, contains multiple copies of one of its peptide products, in this case, the progenitor for TRH, Gln-His-Pro-Gly. Most of the products derived from pro-TRH are targeted into the RSP. Cleavage of the precursor to generate biologically active TRH occurs at paired basic residues by the action of PC1 and PC2 (31, 35, 37) followed by the action of CPE to remove the basic residue(s) (85). Gln-His-Pro-Gly is then amidated by the action of PAM, which uses the C-terminal Gly as the amide donor, and the Gln residue undergoes cyclization to a pGlu residue to yield TRH.

In the last few years, this laboratory has elucidated the processing steps involved in the synthesis of TRH and pro-TRH-derived peptides. An understanding of the biosynthesis and processing of pro-TRH is critical to appreciating how, when, and where modulation of this central regulator of the HPT axis takes place during physiologically appropriate modulation of thyroid function. It is also critical to understanding the function of TRH and other pro-TRH-derived peptides in extrahypothalamic regions of the brain, or outside of the nervous system, as discussed in *Section VI* of this review. Initial studies of TRH biosynthesis were difficult because of the low levels of TRH and other pro-TRH-derived peptides produced in hypothalamic tissue *in vivo*. In the search for a better system to perform these studies, initial work was done by stably transfecting prepro-TRH cDNA into transformed cell lines (86, 87) to attain higher levels of pro-TRH expression. Among the cell lines investigated, only AtT₂₀ (corticotroph, mouse) and RIN 5F (insulinoma, rat) cells were able to efficiently cleave the TRH precursor at paired basic amino acid residues to generate mature TRH, as well as pro-TRH-derived peptides, which were identical to those previously identified *in vivo* (88, 89). GH₄C1 (somatomamotroph, rat) and 3T3 (fibroblast, rat) cells were unable to process the pro-TRH precursor. When these studies were performed, little information was available about PCs, their role in prohormone processing, or their expression in different tissues and cell lines. Retrospectively, it was determined that the cell lines capable of processing proTRH, AtT₂₀ and RIN 5F cells, also express PC1 and PC2 (90).

A model of pro-TRH posttranslational processing was developed in experiments with transfected AtT₂₀ cells expressing rat prepro-TRH (30). By means of Western blot analysis, immunoprecipitation followed by SDS-PAGE, and RIA, it was determined that pro-TRH is present in transfected AtT₂₀ cells and in primary cultures of hypothalamic neurons, an endogenous source of pro-TRH, as a 26-kDa protein (37). Pulse-chase studies indicated that the 26-kDa precursor is cleaved at two mutually exclusive sites to generate the first intermediate forms (Fig. 1). One cleavage generates a 15-kDa N-terminal peptide (prepro-TRH₂₅₋₁₅₁ or 157) and a 10-kDa C-terminal peptide (prepro-TRH₁₅₄ or 160-255). An alternate cleavage generates a 9.5-kDa N-terminal peptide (prepro-TRH₂₅₋₁₀₆ or 112) and a 16.5-kDa C-terminal peptide (prepro-TRH₁₀₉ or 115-255). These cleavage steps occur in the TGN (Fig. 2), before packaging into ISGs (29, 84), in agreement with similar studies of the cellular location of early processing for POMC and somatostatin (SRIF) (53).

In subsequent steps, the 15-kDa N-terminal intermediate moiety of pro-TRH is processed to a 6-kDa peptide, corresponding to prepro-TRH₂₅₋₇₄, and a 3.8-kDa peptide, corresponding to prepro-TRH₇₇₋₁₀₆. It is proposed that processing of the remaining 10-kDa C-terminal fragment produces the 5.4-kDa C-terminal flanking peptide prepro-TRH₂₀₈₋₂₅₅, and the 5.6-kDa peptide prepro-TRH₁₆₀₋₁₉₉ (30, 91). Antisera against prepro-TRH₁₇₈₋₁₉₉ (pFE₂₂) recognizes the 10-, 5.6-, and 2.6-kDa (prepro-TRH₁₇₈₋₁₉₉) C-terminal peptides both in transfected AtT₂₀ cells and in extracts from rat PVN, lateral hypothalamus, and ME. Prepro-TRH₁₇₈₋₁₉₉ is further cleaved to two smaller moieties of 1.6 and 0.84 kDa, prepro-TRH₁₇₈₋₁₈₄ and prepro-TRH₁₈₆₋₁₉₉ (pFQ7 and pSE₁₄), respec-

tively (Fig. 1). Processing of the 9.5-kDa N-terminal fragment arising from the alternative cleavage of the 26-kDa prohormone at residues 107–108 is postulated to result in the production of the N-terminal peptides, prepro-TRH_{25–50}, prepro-TRH_{53–75}, and prepro-TRH_{83–106}, while the 16.5-kDa fragment is processed to produce the 5.6-kDa prepro-TRH_{160–199} and the 5.4-kDa prepro-TRH_{208–255} (Fig. 1).

Recent experiments confirm the proposed model that the C-terminal 10-kDa peptide derived from an initial cleavage at residues Lys¹⁵²-Arg¹⁵³ is the precursor to two peptides of 5.6- (prepro-TRH_{160–199}) and 5.4-kDa (prepro-TRH_{208–255}) (Fig. 1). After the cleavage of basic residues at positions 199–200 or 207–208, the 5.6-kDa peptide is further processed to generate the prepro-TRH_{160–169} and prepro-TRH_{178–184}. prepro-TRH_{178–184} is further cleaved to two novel peptides, prepro-TRH_{178–184} and prepro-TRH_{186–199} (91), Fig. 1).

B. Intracellular sites of pro-TRH processing

Initial studies in 1993 showed that processing of the 26-kDa pro-TRH precursor to smaller intermediates occurred before packaging into ISGs (29). The experimental strategy used to more precisely define the intracellular sites of pro-

TRH processing was to block peptide transport from one cellular compartment to another and to characterize processing that occurred before the point of blockade (84). To study processing that occurs in the RER, peptides were blocked in their exit from this compartment with brefeldin A (BFA) treatment of AtT₂₀ cells expressing the prepro-TRH cDNA (30, 87). BFA is a fungal metabolite that blocks ER-to-Golgi transport of proteins by reversibly inhibiting the exchange of GDP for GTP in the GTP-binding protein that is a key component in the vesicular transport, the ARF protein. This prevents protein recruitment to intracellular membranes and inhibits subsequent vesicle formation. To study the processing steps that occur in the GC, cells were incubated at reduced temperatures. Incubation of cells at 20°C prevents packaging of proteins into ISGs at the TGN and has been used previously to study the processing of other prohormones (53, 92). Figure 2 summarizes the results of these blockade experiments, described more fully below.

When transfected AtT₂₀ cells expressing prepro-TRH were treated with BFA to block ER-to-GC transport, the 26-kDa pro-TRH precursor accumulated 4-fold over control levels, indicating a significant degree of post-ER processing. How-

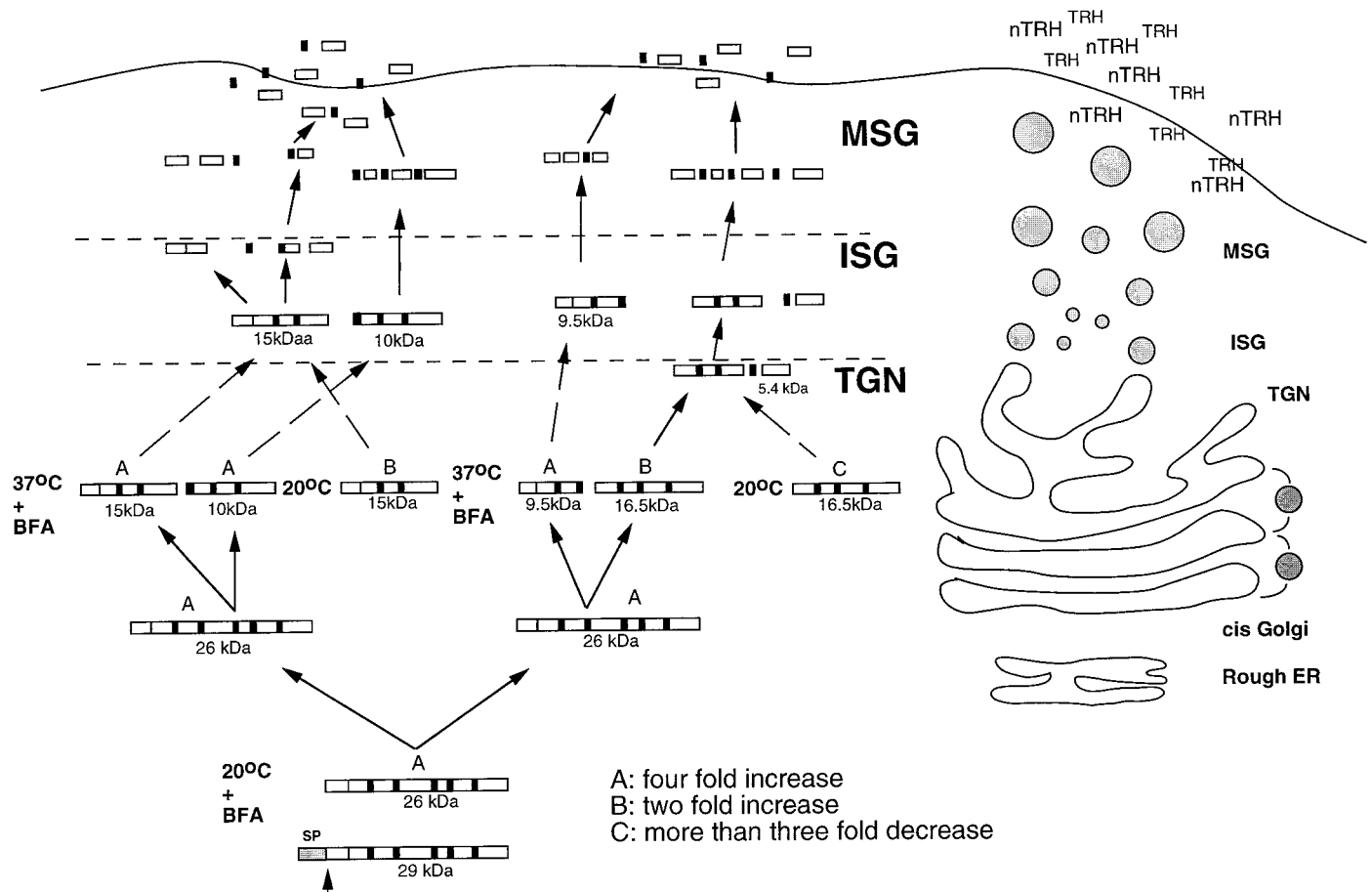


FIG. 2. Schematic representation of the proposed intracellular processing of pro-TRH. On the left is indicated the biochemical processing cascade that is initiated in the GC by PC1 (Section II.B and D). The effect of BFA and temperature blockade on the intracellular trafficking of peptides is described in Section II.B. The PC1 and PC2 enzymatic activities in this cascade are described in Section II.D. On the right, a cartoon showing the distribution of organelles within the RSP. TGN, trans-Golgi-network; MSG, mature secretory granules; ISG, immature secretory granules; nTRH, non-TRH peptides. [Reproduced with permission from I. P. Cruz and E. A. Nillni: *J Biol Chem* 271:22736–22745, 1996 (84).]

ever, some processing of 26-kDa pro-TRH to the 15-kDa and the 9.5-kDa N-terminal intermediates, and the 16.5-kDa and 10-kDa C-terminal intermediates, was seen. The accumulation of the 16.5-kDa intermediate was not as great as that of the 15-kDa intermediate when compared with control levels (4-fold for the 15 kDa and about 2-fold for the 16.5 kDa) (84). To further clarify where the initial site of pro-TRH cleavage occurred, a combination of temperature blockade and BFA treatment was performed. With this strategy it was demonstrated that the 26-kDa precursor protein (accumulated 4-fold) is processed in the GC (possibly in the TGN) to generate the 15-kDa/10-kDa and the 9.5-kDa/16.5-kDa intermediate pairs. When the fate of the 15- and 16-kDa intermediates was analyzed, while they were retained within the TGN at 20 C, the 16-kDa C-terminal intermediate was further processed at basic residues 206–207 to the 5.4-kDa C-terminal peptide. In contrast, the 15-kDa N-terminal intermediate appeared to undergo processing in a post-GC compartment, *i.e.*, the SGs. This observation strongly suggests that these two intermediates follow different paths of processing (84).

Evidence supporting differential distribution for the N- and C-terminal peptides comes from recent immunocytochemical (ICC) studies using transfected AtT₂₀ cells, which indicate that pro-TRH, as well as the 15- and the 6-kDa N-terminal intermediates, are located in the GC and TGN (Fig. 3A). In contrast, end products, including prepro-TRH_{25–50}, prepro-TRH_{160–169}, and TRH, are only present in SGs along the plasma membrane and in cell processes (Fig. 3F). C-terminally directed antisera that recognize pro-TRH and the 16.5- and 5.4-kDa C-terminal peptides result in positive immunostaining in the GC, along the plasma membrane, and in cell processes (Fig. 3H). Thus, C-terminal intermediates appear to reach further along the RSP before processing than their N-terminal counterparts. This differential processing might serve as a mechanism to regulate the timing of production of peptides such as prepro-TRH_{160–169}, prepro-TRH_{178–199}, and prepro-TRH_{53–75}, and possibly TRH. For example, the 16.5-kDa intermediate, which is processed in the TGN, contains prepro-TRH_{178–199} and prepro-TRH_{160–169}. A portion of such peptides, formed before their entry into SGs, might exit the cell via the constitutive pathway to maintain a basal level of release independent of TRH secretion.

Primary cultures of hypothalamic neurons, developed in this laboratory, provided a second model system in which to study pro-TRH processing. After 12–14 days *in vitro* (36), these hypothalamic neurons show growth of neurites similar in morphology to peptidergic neurons. Most neurons are bipolar with long axons containing varicosities, boutons, and growth cones. Many of the growth cones are in contact with neurites of other neurons. Dendrite-like structures are also observed. N-terminal antiserum in these cultured hypothalamic neurons (36) stains the GC, ISGs budding from the GC, and terminal boutons (Fig. 3, B and E). Immunoelectron microscopy (IEM) confirms positive staining in the GC (Fig. 3E). Some ISGs budding from the TGN-like were also stained. Since positive staining was detected in boutons, it was proposed that these intermediates are processed to mature pro-TRH-derived peptides near the axon terminal, before secretion. ICC using anti-TRH, anti-prepro-TRH_{53–75}, and anti-prepro-TRH_{160–169} showed positive staining only in

the neuronal processes and axon terminals while the soma remained unstained (Fig. 3G). The positive staining observed in neuronal processes and axon terminals with these antibodies was confirmed by IEM (Fig. 3G, *inset*).

Note that in transfected AtT₂₀ cells using the same antiserum, immunostaining was seen only in GC and ISGs, but not in cellular processes (Fig. 3A). ICC on transfected AtT₂₀ cells using anti-TGN₃₈, a TGN marker, indicated that the TGN can be projected away from the GC toward budding ISGs (Fig. 3C). Thus, in AtT₂₀ cells processing of the 15-kDa N-terminal peptide takes place somewhere between the TGN and ISGs, suggesting that prepro-TRH_{25–50}, prepro-TRH_{53–75}, and prepro-TRH_{83–106} are already formed by the time SGs mature.

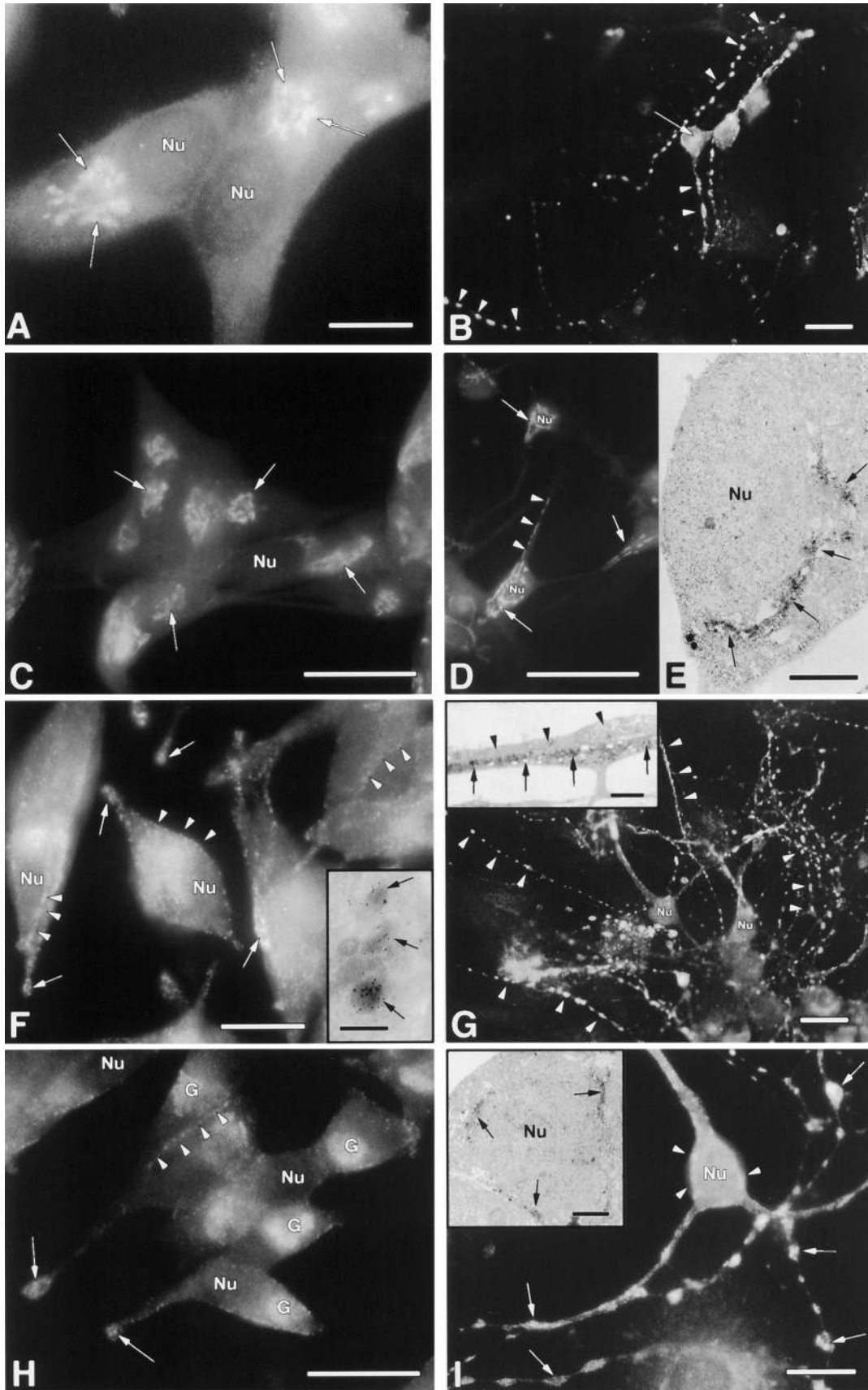
Using the C-terminal antiserum that recognizes pro-TRH and the 16.5-, 10-, and 5.4-kDa peptides, positive staining was visualized in a patchy cytoplasmic distribution (Fig. 3I), often closely associated with the nucleus. Immunoreactivity was also observed in neuronal boutons, axon terminals, and unbranched growth cones. IEM confirmed that the patchy cytoplasmic areas were within the ER and GC (36) (Fig. 3I, *inset*); all layers of the GC (*cis*, medial, and *trans*) were immunostained with this antibody (36). The contrasting staining patterns for the two antisera (N- and C-terminal) (Fig. 3, B and I) suggest the existence of a different peptide distribution for N-terminal *vs.* C-terminal peptides, possibly due to different intracellular routing of intermediates to SGs. However, no conclusive data are yet available to confirm this hypothesis.

C. Tissue-specific processing of pro-TRH

Increased knowledge of differential processing has led to a better understanding of how multiple biological peptides, with different functions, are generated from the same prohormone. This concept is further reinforced by the observation that certain regions in the brain can give rise to several different pro-TRH-derived peptides in addition to, or instead of, TRH. This will become clear in a review of the neuroanatomical distribution of pro-TRH and pro-TRH-derived peptides.

TRH-positive axons in the ME originate from neuronal perikarya in the PVN. The PVN is composed of two major components, the magnocellular and the parvocellular divisions. The parvocellular division contains most of the TRH neurons that project to the ME. A large population of immunoreactive neurons is located in medial and periventricular parvocellular subdivisions, organized in a triangular configuration, symmetric to the dorsal aspect of the third ventricle, whereas the anterior parvocellular subdivision neurons are more dispersed (93). However, not all TRH-containing neurons in the PVN project to the ME (94). In addition to the PVN, TRH neurons are present in many other regions of the hypothalamus (93). As these populations of neurons have no known projections to the ME, and are not regulated in conjunction with the thyrotropic neurons of the PVN, it is presumed that they do not subservise a direct hypophysiotropic function.

The largest concentration of hypothalamic TRH neurons outside of the PVN are found in the dorsomedial nucleus,



lateral hypothalamus, and preoptic area, including medial, periventricular, suprachiasmatic, and the sexual dimorphic nucleus of the preoptic area (93). In addition to the PVN, TRH neurons are present in many other regions of the CNS including regions in the diencephalon, telencephalon, mesencephalon, myelencephalon, and spinal cord (Table 1). An extensive anatomic description of TRH neurons and TRH fibers in these tissues has been reported previously (93). While the presence of TRH in these regions is clear, relative levels are largely unknown because most mapping has depended on ICC detection in animals pretreated with colchicine. Detailed microdissection studies, combined with RIA, have been done for some areas, including the brainstem and hippocampus. These studies do not require colchicine pretreatment, are more quantitative than ICC detection, and reveal an even broader distribution for TRH than appreciated by ICC. However, their neuroanatomical resolution falls short of that of ICC, and these latter studies are the primary source of TRH data presented in Table 1.

In several areas of the brain where production of pro-TRH is found, TRH and pro-TRH-derived peptides are also detected (Table 1). In the case of pro-TRH, the reticular nucleus of the thalamus contains abundant prepro-TRH mRNA and several pro-TRH-derived peptides in their extended forms, but does not contain mature TRH (95). Moreover, the N-terminal extended forms of TRH, TRH-prepro-TRH₁₆₀₋₁₆₉ and TRH-prepro-TRH₁₇₈₋₁₉₉ are major end products of pro-TRH processing in the olfactory lobe (OB) (96, 97), but not in the hypothalamus where pro-TRH is completely processed to non-TRH peptides and TRH (96). In the ME, PVN, and preoptic area (POA), pro-TRH is fully processed to its mature forms, while in the OB less than 60% of N-terminal prepro-TRH₂₅₋₅₀ is formed. Similarly, while in the OB the 10-kDa C-terminal intermediate (see Fig. 1) is the main end product, in the ME and PVN this intermediate is fully processed to prepro-TRH₁₆₀₋₁₆₉ and prepro-TRH₁₇₈₋₁₉₉ (Fig. 1). Finally, in the POA partial processing of the 10-kDa peptide is observed, and the lateral hypothalamus contains lower levels of both N- and C-terminal pro-TRH-derived peptides as compared

with the ME and POA (E. A. Nillni, unpublished data). Table 1 summarizes the current information regarding neuroanatomical distribution of intermediate forms of pro-TRH processing, mature pro-TRH-derived peptides, and TRH peptides in perikarya and fibers.

D. The role of PCs and CPs

The PCs are a family of seven subtilisin/kexin-like endoproteases including furin, PC1 (also known as PC3), PC2, PC4, PACE4, PC5-A (also known as PC6-A), its isoform PC5-B (also known as PC6-B), PC7 (also known as LPC), and PC8 (also known as SPC7) (32–34, 98–101). The structure of these serine proteinases resembles both the bacterial subtilisins and yeast kexin (90, 102, 103). These enzymes cleave at the C-terminal side of single, paired, or tetra basic amino acid residue motifs (104), followed by removal of remaining basic residue(s) by CPs (50, 51). The selective expression of PC1 and PC2 in endocrine and neuroendocrine cells suggests they are significant in prohormone processing (32, 90, 98, 105). PC1 and PC2 have been shown to process pro-TRH (31, 35, 37, 106), proinsulin (104, 107, 108), proenkephalin (109), prosomatostatin (110, 111), and POMC (112, 113) to various intermediates and end products in coexpression experiments.

Like their substrates, PC1 and PC2 undergo maturation from larger precursor proteins. Maturation of pro-PC1 begins in the ER and continues in the TGN (90). In contrast, pro-PC2 maturation begins in the TGN and continues in the SGs; active PC2 and PC1 accumulate in SGs (114, 115). PC2 has been proposed to produce smaller peptides from intermediates resulting from cleavage by PC1 (109, 112, 113). Consistent with this, PC1 is implicated in the early cleavage of POMC (112, 113, 116) and proinsulin (117). Thus, the compartments where these enzymes are active also are determined by differential timing of their respective maturation.

The distribution of neurons containing mRNAs encoding pro-TRH, PC1, and PC2 in the PVN and other areas of the

FIG. 3. Subcellular localization of pro-TRH, intermediate and end products of processing, and TRH in transfected AtT₂₀ cells encoding prepro-TRH, and in primary cultures of hypothalamic neurons. The cells were fixed with 4% paraformaldehyde followed by immunostaining with different antibodies against the pro-TRH sequence. Fluorescein isothiocyanate conjugated to goat antirabbit globulin was used as a probe. Panel A, AtT₂₀ cells: positive staining in the GC and TGN (arrows) using an antibody against pro-TRH and N-terminal intermediate forms (anti-pCC₁₀). Bar = 25 μ m. Panel C, AtT₂₀ cells: cells immunostained with anti-TGN38 (arrows), a TGN marker. Bar = 50 μ m. Panel F, AtT₂₀ cells: a typical positive staining along the plasma membrane (arrow heads), a common granule distribution of corticotropic cells, and processes (arrows) using anti-non-TRH peptides and anti-TRH. Bar = 25 μ m. Panel F, AtT₂₀ cells; inset: typical positive staining of SGs by IEM using anti-pST₁₀ antibodies (5 nm gold particles). Bar = 200 nm. Panel H, AtT₂₀ cells: positive staining in the GC and processes using an antibody that recognizes pro-TRH and C-terminal intermediate forms (anti-pYE₁₇). Bar = 50 μ m. Panel B, Hypothalamic neurons: positive staining in the GC (arrow) and TGN using an antibody against pro-TRH and N-terminal intermediate forms, and in all boutons distributed along the neuronal processes (arrowheads) Bar = 50 μ m. Panel D, hypothalamic neurons: cells immunostained with anti-TGN38, a TGN marker. Bar = 50 μ m. Panel E, hypothalamic neurons, a higher magnification of panel B showing positive staining in stacked Golgi cisternae and in some forming granules (arrows) using the peroxidase-DAB reaction (arrows) Bar = 5 μ m. Panel G, hypothalamic neurons: positive fluorescence is observed only in neurites (arrowheads) and axon terminals, while the cell body remain unstained. Bar = 50 μ m. Panel G, hypothalamic neurons; inset: an IEM of neurites using peroxidase-DAB staining reaction. Of the two adjacent neurites shown, the lower one is positively stained (large arrows), whereas the upper one is negative (small arrow). Bar = 1 μ m. Panel I, hypothalamic neurons: positive staining in several areas of the cell body (arrowheads) and in all boutons distributed along the neuronal processes (arrowheads) using an antibody against pro-TRH and C-terminal intermediate forms. Bar = 25 μ m. Panel I, hypothalamic neurons, inset: a higher magnification of cytoplasmic areas from panel I showing positive staining in the endoplasmic reticulum and GC (arrows) as well as in SG near the plasma membrane (arrows). Bar = 2 μ m. nu, Nucleus; G, Golgi complex. The polyclonal antibodies used in this ICC are as follow: Anti-pCC₁₀ [made against a synthetic decapeptide (Cys-Lys-Arg-Gln-His-Pro-Gly-Lys-Arg-Cys)], which recognizes prepro-TRH₂₅₋₂₅₅ (26 kDa) prepro-TRH₂₅₋₁₅₁ (15 kDa) prepro-TRH₂₅₋₁₁₂ (9.5 kDa) prepro-TRH₂₅₋₇₄ (6 kDa). Anti-pYE₁₇ (made against prepro-TRH₂₄₀₋₂₅₅), which recognizes prepro-TRH₂₅₋₂₅₅ (26 kDa) prepro-TRH₁₁₅₋₂₅₅ (16.5 kDa), prepro-TRH₁₆₀₋₂₅₅ (10 kDa), prepro-TRH₂₀₈₋₂₅₅ (5.4 kDa), anti-pST₁₀ (made against preproTRH₁₆₀₋₁₆₉), and anti-TRH. [Panels B, D, G, and I were reproduced with permission from E. A. Nillni *et al.*: *Endocrinology* 137:5651–5661, 1996 (36). © The Endocrine Society.]

TABLE 1. Current information available regarding pro-TRH biosynthesis, processing in different tissue location, its relation to PC1 and PC2, neuroendocrine inputs, and physiological significance

Tissue	pro-TRH-derived peptides	TRH		PCs			Coexpressing PCs and pro-TRH	Neuroendocrine inputs to TRH neurons	Physiological significance
		Perikarye	Fibers	PC1	PC2	PC5			
Telencephalon									
Olfactory bulb	Partial processing prepro-TRH154-169 prepro-TRH172-199								
Accessory			+						
External plexiform layer		+		+++	++	++		PC2	
Internal plexiform layer		+						PC2	
Mitral cells				+	++	++++			
Ependymal cells				0	0	0			
Granule cells				0/+	+	0			
Periglomerular cells				+	++	++			
Glomerular layer		+	+					PC2	
Anterior olfactory nucleus		+		+++	++++	++++			
Tenia tecta				++++	+++	0		PC1	
Hippocampal formation									
CA1				+++	+++++	++			
CA2		+		++	++++	+			
CA3		+		++++	+++++	+++++			
Scattered hilar cells				++++	+++	++++			
Stratum oriens			+						
Dentate gyrus									
Molecular Layer		+		0	0	0		PC1,PC2	
Granular Layer		+		+++++	+++	+		PC1,PC2	
Nucleus accumbens			+	+	++	0			
Caudate-putamen		+	+	+++++	+++	0		PC1,PC2	
Amygdala									
Medial nucleus		+	+	+++	++	+++		PC1,PC2	
Central nucleus		+	+	0/+	0/+	+		PC1,PC2	
Anterior cortical nucleus				++	+	+++		PC1,PC2	
Basolateral nucleus				+++	++++	+++		PC1,PC2	
Septum			+	++	++++	++			
Bed nucleus, anterior commissure		+							
Bed nucleus, stria terminalis		+	+	+++	++	+		PC1,PC2	
Diagonal band of Broca		+	+					PC1,PC2	
Entorhinal cortex		+							
Medial preoptic area				+++	+	++		PC1,PC2	
Pyramidal cortex			+						
Diencephalon									
Thalamus									
Anteroventral nucleus				+++	++	0			
Anteromedial nucleus				++	++++	0/+			
Anterodorsal nucleus				+++++	++++	0/+			
Centrolateral nucleus				++	++++	0/+			
Centromedial nucleus				+++	++++	+++++			
Mediodorsal nucleus				+++	++++	0			
Paraventricular nucleus			+	+++	++++	+++++			
Paratenial nucleus			+						
Reticular nucleus	Some intermediates, no TRH		+					0	
Reuniens nucleus				0	+++	0/+			
Rhomboid nucleus				+++	++++	+++			
Ventrolateral nucleus				0/+	++++	0			
Ventromedial nucleus			+	0/+	+++	0			
Dorsal lateral geniculate nucleus				0	++++	0			
Medial geniculate nucleus									
Dorsal				0/+	++++	0			
Ventral				0/+	+++	0			
Medial				0/+	++++	0			
Marginal Zone				+++	+	0			
Habenua									
Medial									
Lateral part				++	+	0			
Medial part				+	++++	0			
Lateral			+	++	+	0			
Hypothalamus									
Anterior hypothalamus		+	+						
Suprachiasmatic nucleus				+++	++	++		PC1,PC2	
Periventricular nucleus				++++	+	+++		PC1,PC2	
Supraoptic nucleus		+		+++++	+++	+++++		PC1,PC2	

TABLE 1. Continued

Tissue	pro-TRH-derived peptides	TRH		PCs			Coexpressing PCs and pro-TRH	Neuroendocrine inputs to TRH neurons	Physiological significance
		Perikarye	Fibers	PC1	PC2	PC5			
Paraventricular nucleus	pro-TRH is fully processed							NE+, E-?, DA+, 5HT+, NPY-, Som-, EOPs-, VIP+?, GABA-?, IL-1-	Theromoregulation, T ₃ /T ₄ feedback, stress, TSH circadian rhythm, starvation, response to infection. pFE22 and pSE14 increased during suckling
Magnocellular part		+	+	+++++	++	+++++	PC1,PC2		
Parvocellular part		+	+	++	+++	+++	PC1,PC2		
Arcuate nucleus		+	+	+++	++	++	PC1,PC2		
Lateral hypothalamic area	Lesser processing as compared with PVN and ME	+	+	+++++	+++++	+++++	PC2		
Median eminence	TRH + non-TRH peptides		+					NE+, E?, DA-, Som-, EOPs-, IL-1?	Theromoregulation. T ₃ /T ₄ feedback, stress, TSH circadian rhythm, starvation, response to infection
Ventromedial nucleus			+	+	++	+	PC2		
Dorsomedial nucleus		+	+	+++	+++	+++	PC1,PC2		
Medial mammillary nucleus				+	++++	+++			
Lateral mammillary nucleus				+	+++	+			
Perifornical region		+							
Premammillary nucleus			+						
Preoptic area	pro-TRH is fully processed	+	+						
Subfornical organ			+						
Mesencephalon and metencephalon									
Locus coeruleus			+	++	++++	+++			
Motor nucleus V			+						
Oculomotor nucleus			+	+++	++++	+++			
Parabrachial nucleus			+						
Periaqueductal gray	pro-TRH is fully processed	+	+	++	+++	++	PC1,PC2		Possible pain modulation
Raphe magnus nucleus				+++	+	0	PC1,PC2		
Red nucleus			+	++	++++	++++			
Substantia nigra, pars lateralis		+							
Substantia nigra, pars compacta				+++	+++	+++			
Substantia nigra, pars reticulata				+	+++	+++			
Trochlear nucleus			+						
Myelencephalon									
Cochlear nucleus		+		++++	+++	++			
Dorsal motor nucleus, vagus		+	+						
External cuneate nucleus		+							
Facial nucleus			+	+++	++++	++++			
Hypoglossal nucleus			+	+++	+	++++			
Inferior olive			+	++	++++	++++			
Lateral reticular nucleus		+							
Nucleus ambiguus			+						
Nucleus tractus solitarius									
Medial			+	++++	++	++++			
Lateral				++	+++	++++			
Pontine nuclei		+		+++	++++	++++	PC1,PC2		
Raphe nuclei									
Dorsal		+	+	+++	+++	++++	PC1,PC2		
Median		+	+	+++	++	+++	PC1,PC2		
Reticular formation			+						
Spinal Cord	pST10 detected								
Anterior funiculus			+						
Central canal			+						
Dorsal horn (laminae II and III)		+	+						
Intermediolateral column			+						
Lateral funiculus			+						
Ventral horn (lamina IX)			+						
Cell lines									
Transfected AtT20 cells	pro-TRH is fully processed			PC1	low PC2		PC1,PC2		
Transfected RIN 5F cells	pro-TRH is fully processed			PC1	PC2				
Transfected GH4C1 cells	no pro-TRH processing								
Primary hypothalamic neurons	pro-TRH is fully processed			PC1	PC2		PC1,PC2		

The major source of contribution to this table comes from literature cited in this review and from our published and unpublished results.

brain has been determined using *in situ* hybridization (106) (Table 1). The glomerular layer of the OB displays coexpression of prepro-TRH with PC2 mRNA, but not PC1, whereas in the tenia tecta coexpression of mRNA for prepro-TRH with PC1, but not PC2, is evident. The PVN displays prepro-TRH mRNA coexpression with both enzymes, whereas the lateral hypothalamus shows coexpression of prepro-TRH mRNA with PC2 mRNA, but not PC1 mRNA. Double *in situ* hybridization indicates that in the PVN, PC2 mRNA is present in 60–70% of TRH neurons, and PC1 is present in 37–46% of TRH neurons (118). Even though these investigators found a trend for more coexpression of mRNA for prepro-TRH with PC2 than PC1, coexpression alone does not define which enzyme is more important in the processing of pro-TRH *in vivo*.

During the last few years, Nillni's laboratory has provided unequivocal evidence for the role of PC1 and PC2 in the processing of pro-TRH (31, 35–37). In coinfection experiments, where recombinant vaccinia viruses are used to coexpress PC1, PC2, PACE4, PC5-B, and furin, together with rat prepro-TRH in constitutively-secreting LoVo cells or in the regulated-secreting endocrine GH₄C1 cell line, RIA of LoVo-derived secreted products demonstrates that furin cleaves the precursor to generate both N- and C-terminal intermediates, while PC5-B does not produce any peptide. PC1, PC2, and PACE4 only produce N-terminal intermediates, and less efficiently than furin. Interestingly, in LoVo cells, furin cotransfection produces TRH-Gly at much greater levels than any of the other PC enzymes. Recent data indicate that furin, which is ubiquitously expressed in all tissues, may serve a role in processing of prosomatostatin within the constitutive pathway (110, 111). Since LoVo cells only contain the constitutive secretory pathway, these results suggest that pro-TRH can be processed to a certain extent without entry into the RSP. However, caution must be taken with this interpretation because under conditions of viral expression, the unusually high level of virus in coinfecting cells can produce disruption of cellular compartments. The products resulting from coexpression of prepro-TRH with either furin or PC1 are similar, in agreement with their similar specificity observed in a number of cell coexpression experiments (109, 111) and *in vitro* data (119).

In GH₄C1 cells, PC1, PC2, furin, PC5-B, and PACE4 produce both N-terminal and C-terminal peptides derived from pro-TRH. Significantly, TRH-Gly and TRH are produced in highest amounts by PC1, PC2, and furin. Further analysis of the cleavage specificity of PC1 and PC2 reveals that PC1 is primarily responsible for cleavage of the entire TRH precursor to mature TRH, as it can generate all products at significantly higher levels than PC2 (Fig. 4). While 7B2 is known to be involved in the maturation of PC2 (Fig. 4) (90, 120) it does not augment the ability of PC2 to cleave pro-TRH to either N- or C-terminal forms. Subsequently, we have examined the role of PC1 and PC2 in the formation of prepro-TRH_{178–199} by coexpressing rat prepro-TRH cDNA with PC1, PC2, and 7B2 in GH₄C1 cells (91). PC1 effectively cleaved pro-TRH to immunoreactive forms recognized by anti-prepro-TRH_{178–199}, while PC2 played a minor role, even in the presence of 7B2 (Fig. 1).

Even though PC1 displays a greater ability to process

pro-TRH than PC2, PC2 can process certain regions of the pro-TRH sequence (37). For example, in cells coexpressing pro-TRH and PC2, but not pro-TRH and PC1, greater quantities of 2.8-kDa peptide (pEH₂₄, preproTRH_{83–106}) relative to 3.8-kDa peptide (TRH-pEH₂₄, prepro-TRH_{77–106}) are found, while the converse is true for PC1 coexpressing cells. These data suggest that PC2 may be important in generating TRH from this intermediate. In support of a physiological role for both PC1 and PC2, pro-TRH is coexpressed and colocalized with PC1 and PC2 in primary cultures of hypothalamic neurons (Fig. 5) (37).

SDS-PAGE fractionation reveals that PC2 has a cleavage specificity that differs from that of PC1. In cells coinfecting with PC1 and pro-TRH, two prominent moieties of 5.6 kDa and 2.6 kDa are formed. The latter is similar in size to prepro-TRH_{178–199}. In cells coinfecting with PC2 and proTRH, the 5.6-kDa peptide is not present, but the 2.6-kDa peptide and a smaller form of about 1.6 kDa are observed. Microsequencing analysis of prepro-TRH_{178–199} peptide incubated *in vitro* with purified PC2 demonstrates one specific cleavage at Arg¹⁸⁵ (. . . Glu¹⁸²-Leu¹⁸³-Gln¹⁸⁴-Arg¹⁸⁵-↓-Ser¹⁸⁶-Trp¹⁸⁷-Glu¹⁸⁸-Glu¹⁸⁸-Lys¹⁸⁹. . .) generating two novel peptides, pFQ₇ and pSE₁₄ (Fig. 1).

These two novel peptides, prepro-TRH_{178–184} and prepro-TRH_{186–199} are present in rat PVN, lateral hypothalamus, and ME (91). Thus, the antibody generated against the prepro-TRH_{178–199} sequence recognizes the 10-kDa peptide, a 5.6-kDa form that probably is prepro-TRH_{160–199} (30), a 2.6-kDa peptide that is prepro-TRH_{178–199}, and two smaller moieties of 1.6 and 0.84 kDa that are proposed to be prepro-TRH_{186–199} and prepro-TRH_{178–184}, respectively (Fig. 1). Figure 6 shows a diagrammatic representation of rat pro-TRH and its cleavage by PC1 and PC2 as proposed from the most recent studies and compared with previous *in vitro* studies (31, 35, 91).

Table 1 combines current neuroanatomical distribution data for prepro-TRH mRNA, pro-TRH, and pro-TRH-derived products with the distribution of PC mRNAs and enzymes. Also included is a summary of pro-TRH processing in transfected cells and primary cultures of hypothalamic neurons and pituitary cells. Several important conclusions can be drawn. pro-TRH is widely distributed in the hypophysiotropic and extrahypophysiotropic areas of the brain. The widespread expression of pro-TRH, PC1, and PC2 mRNAs, with their overlapping distribution in many areas of the rat CNS, indicates the striking versatility provided by tissue-specific processing in generating quantitative and qualitative differences in non-TRH peptide products as well as TRH. Examples of these differences for several tissues are presented in the first column of the table and described above. A most striking example is the reticular nucleus of the thalamus, where PC1 and PC2 are not coexpressed with pro-TRH. TRH is not produced in this nucleus, indicating a central role for PC1 and PC2 in maturation to TRH. However, other pro-TRH intermediates are present in the reticular nucleus, suggesting PCs other than PC1 and PC2 might be involved in processing of pro-TRH for this particular region of the CNS (106).

Differential processing has been reported for other pro-hormones, and these differences relate to alterations in the expression of various PCs within different cell types. POMC

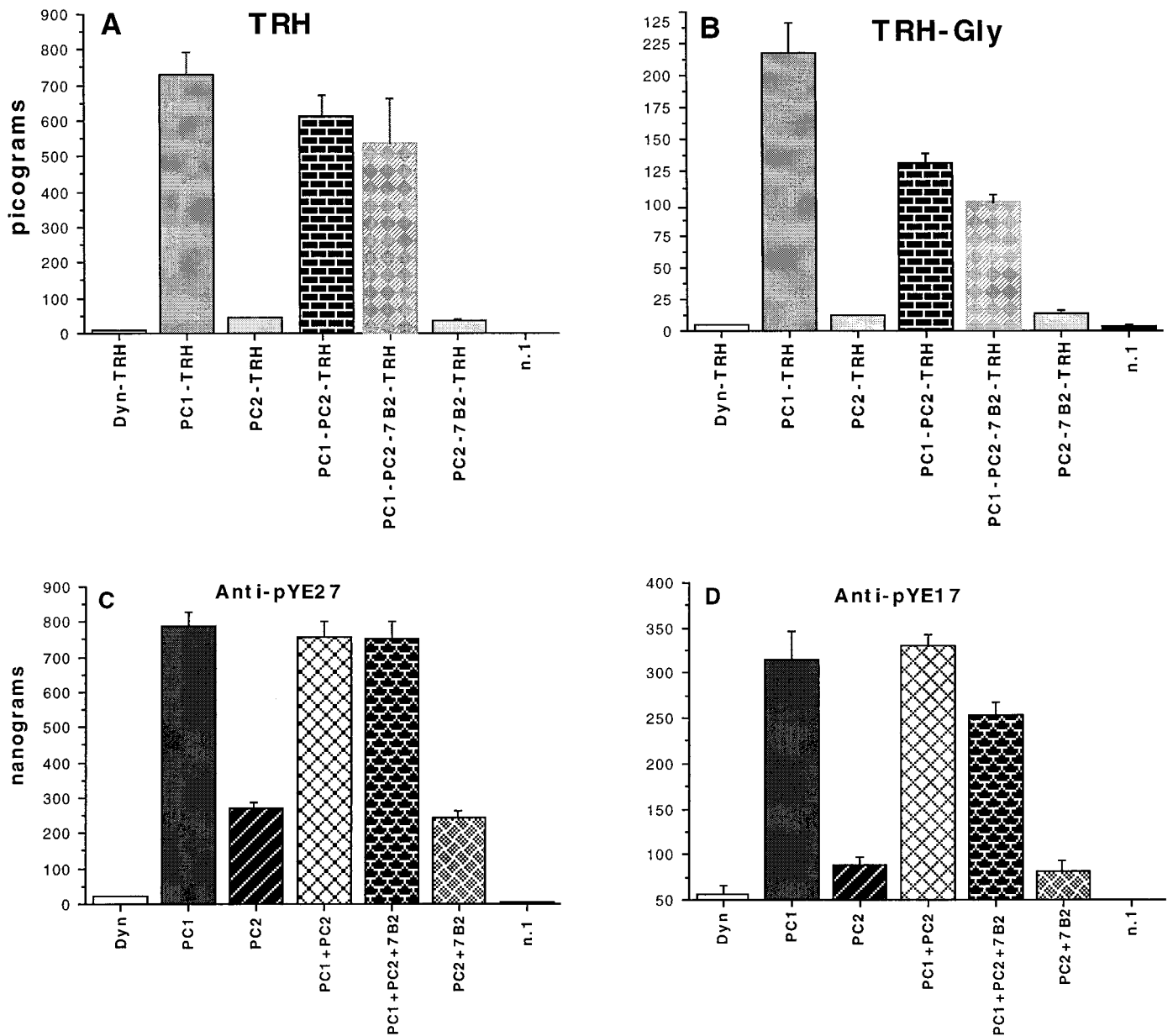


FIG. 4. Cleavage of pro-TRH as determined using anti-TRH (A) RIA, anti-TRH-Gly (B), using anti-pYE₂₇ (C) RIA, and anti-pYE₁₇ (D) RIA. Coinfections of pro-TRH were done with Dynorphin as a control, PC1, PC2, PC1-PC2, PC1-PC2-7B2, and PC2-7B2 in GH4C1 cells. n, Noninfected cells. RIAs were performed against resuspended serum free media. Cell means of recognized products in picograms are plotted against the indicated coinfecting construct. Data are the mean values of six identical wells per condition, with $P < 0.05$ on Tukey-Kramer. [Reproduced with permission from P. Schaner *et al.*: *J Biol Chem* 272:19958–19968, 1997 (37).]

is processed primarily to ACTH, β -endorphin, and N-POMC₁₋₇₇ in the anterior pituitary (melanotrophs). In turn, these products are further processed to α -MSH, β -endorphin₁₋₃₁, N-POMC₁₋₄₉, and γ -MSH in the intermediate lobe and brain (38). Differential processing of a common polypeptide precursor is dependent upon the processing enzymes expressed in each specific cell type. Proenkephalin, which contains seven identical copies of met-enkephalin, is processed to large intermediate forms in the adrenal medulla, whereas this precursor is cleaved primarily to the pentapeptide met-enkephalin in the brain (39). The biological actions of substance P (SP) depend on the enzymatic pro-

cessing of its precursor by the processing enzymes prolylendopeptidase to yield SP₅₋₁₁, and endopeptidase 3.4.24.11 to yield SP₁₋₇. While SP₁₋₇ acts as an analgesic, inhibits aggression, and enhances learning and memory, the SP₅₋₁₁ enhances pain transmission, stimulates aggression, and blocks learning and memory (40). In the brain, procholecystinin (pro-CCK) is processed to produce only CCK8 amide, while in the gut the precursor is cleaved to larger molecules, such as CCK12, 22, 33, 38, 58, and 83 amide (121). Transfection experiments have shown that proneuropeptide Y (pro-NPY) can be cleaved by cell lines expressing either PC1 or PC2, but pro-NPY is primarily processed by PC2 in superior cervical

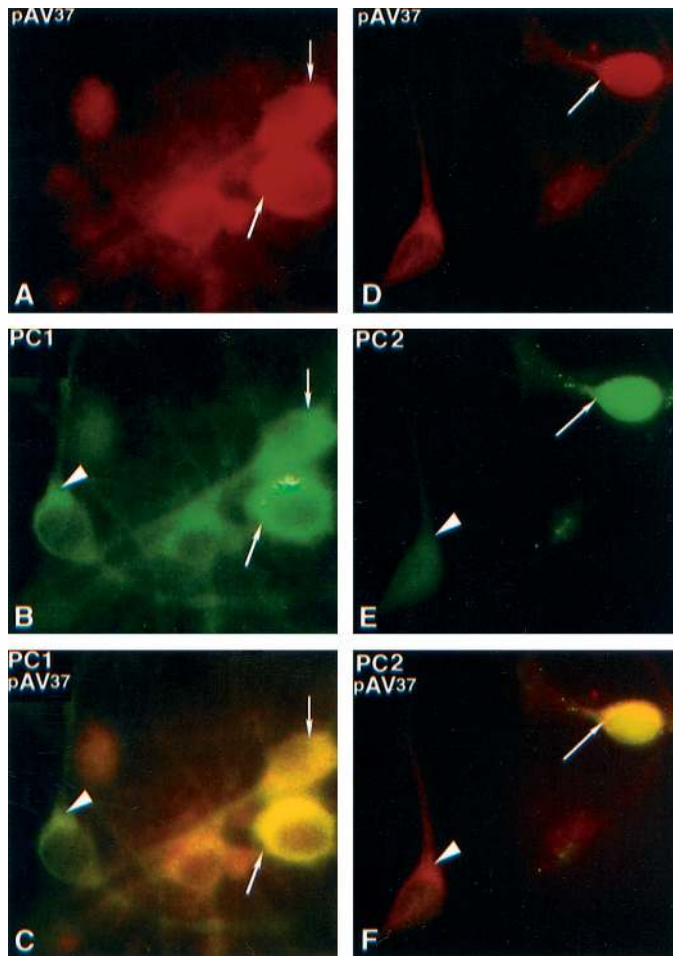


FIG. 5. Colocalization of pro-TRH with PC1 and PC2. Neuronal cells cultured for up to 14 days in four-chamber labTeck slides were fixed with 4% paraformaldehyde followed by immunoreaction with anti-PC1 or anti-PC2. Fluorescein isothiocyanate conjugated to goat anti-rabbit globulin was used as a probe. Texas Red-X-succinidylester directly conjugated to anti-pAV₃₇ antibodies was used as a probe for pro-TRH. Panel A shows positive immunostaining for pro-TRH peptides (red color, arrows). Panel B shows positive immunostaining for PC1 (green color, arrow and arrowhead). Panel C shows the protein colocalization of pro-TRH and PC1 (yellow-orange color indicated by arrows). Some neuronal cells contain PC1 but not pro-TRH (panels B and C, arrowhead). Panel D shows positive immunostaining for pro-TRH peptides (red color, arrow). Panel E shows positive immunostaining for PC2 (green color, arrow and arrowhead). Panel F shows the protein colocalization of pro-TRH and PC2 (yellow-orange color indicated by arrow and arrowhead). Some neuronal cells contain PC1 but not pro-TRH (panels B and C, arrowhead). Thirty-millimeter slides were digitized with a video camera and appropriate macro lens using BioVisionframe grabber software (Perceptics Corp., Knoxville, TN). Images of the red and green planes were combined using Adobe Systems, Mountain View, CA) to show areas of colocalization. The resulting images were printed with a Mitsubishi CP210 dye sublimation printer (Apunix Computer Services, San Diego, CA). [Reproduced with permission from P. Schaner *et al.*: *J Biol Chem* 272: 19958–19968, 1997 (37).]

ganglia (122). Thus, differential processing of neuropeptides including pro-TRH, pro-NPY, POMC, pro-CCK, SP, and proenkephalin provides a critical mechanism through which cells regulate the levels of specific peptides to fulfill different physiological requirements, a mechanism potentially more versatile than the alternative splicing of mRNA.

As mentioned in Section IIA, CPs remove remaining C-terminal basic residues from prohormone intermediates that are initially cleaved by PCs. Experiments with the *fat/fat* mouse model of CPE deficiency (123) support a role for CPE in the processing of pro-TRH (85). Mice homozygous for the *fat/fat* mutation are obese, diabetic, and infertile. These mice have a missense (Ser to Pro) mutation at CPE residue 202 that abolishes enzymatic activity (123). Hypothalamic TRH levels are depressed 65% in *fat/fat* mice relative to heterozygous controls. SDS-PAGE demonstrates hypothalami from both *wt/fat* and *fat/fat* mice contain moieties different from those of the *wt/wt* mice. Specifically, 6.5-, 3.0-, 2.6-, and 1.6-kDa forms of the pro-TRH sequence are detected, and their levels differ significantly between the two groups. Compared with *wt/fat* mice, *fat/fat* mice hypothalami contain 20-, 3-, and 2-fold elevations in the 6.5-, 3.0-, and 2.6-kDa species. These data indicate that the *fat/fat* mutation produces qualitative changes in pro-TRH processing, and that CPE is involved in the later stages of pro-TRH processing. However, cell transfection experiments would help to rule out secondary phenotypic changes caused by the CPE mutation in these mice. Further, since hypothalami from *fat/fat* mice contain immunoreactive TRH, additional CPs must also be able to process pro-TRH to TRH, assuming that the TRH detected is not a cross-reactive non-TRH species. CPs such as carboxypeptidase D, with similar enzymatic properties to CPE, are also present in compartments of the secretory pathway and are distributed in many tissues, including the brain (50).

Interestingly, in *fat/fat* mice, levels of TSH, T₃, and T₄ were normal, suggesting that 34% of normal TRH levels is sufficient to maintain the thyroid function. This last observation is important because it is hypothesized that the five identical progenitor sequences of TRH contained in the prohormone may not be processed to mature TRH at all times, and that only a few of them may be needed to maintain the thyroid function. In cultured hypothalamic cells at steady state, previous studies had shown that the ratio of mature TRH to prepro-TRH_{25–50} and the 5.4-kDa C-terminal peptide (Fig. 1) is 3:1 instead of the theoretical 5:1 for complete processing (36). A similar ratio is seen in transfected AtT₂₀ cells (30). In the rat brain, the ratio of TRH to other pro-TRH-derived peptides sequences is almost 1:1 in the hypothalamus, and 1.5:1 in the olfactory lobe (89). The above ratios are indicative of incomplete yields of TRH from pro-TRH, although only if degradation rates do not contribute significantly to these various ratios. Further, the relative immunoreactivities of various antisera used to their iodinated tracers has not been defined, leaving exact molar ratios difficult to calculate. Still the range of TRH to pro-TRH-derived peptides found in various tissues makes it likely that full TRH yields are not achieved in all, or even most, tissues.

If not all TRH progenitor sequences are cleaved from pro-TRH, the resulting TRH progenitor sequences, linked to amino- or carboxyl-terminal extensions, would not be detected in usual TRH RIAs. The TRH assay is specific for mature TRH, needing both the amino-terminal pyro-Glu and carboxyl-terminal Pro-amide for detection. Nonimmunoreactive TRH progenitors might also retain untrimmed basic amino acid residues, carboxyl-terminal glycine residues, etc. Another explanation for the lower than 5:1 ratio of TRH to other

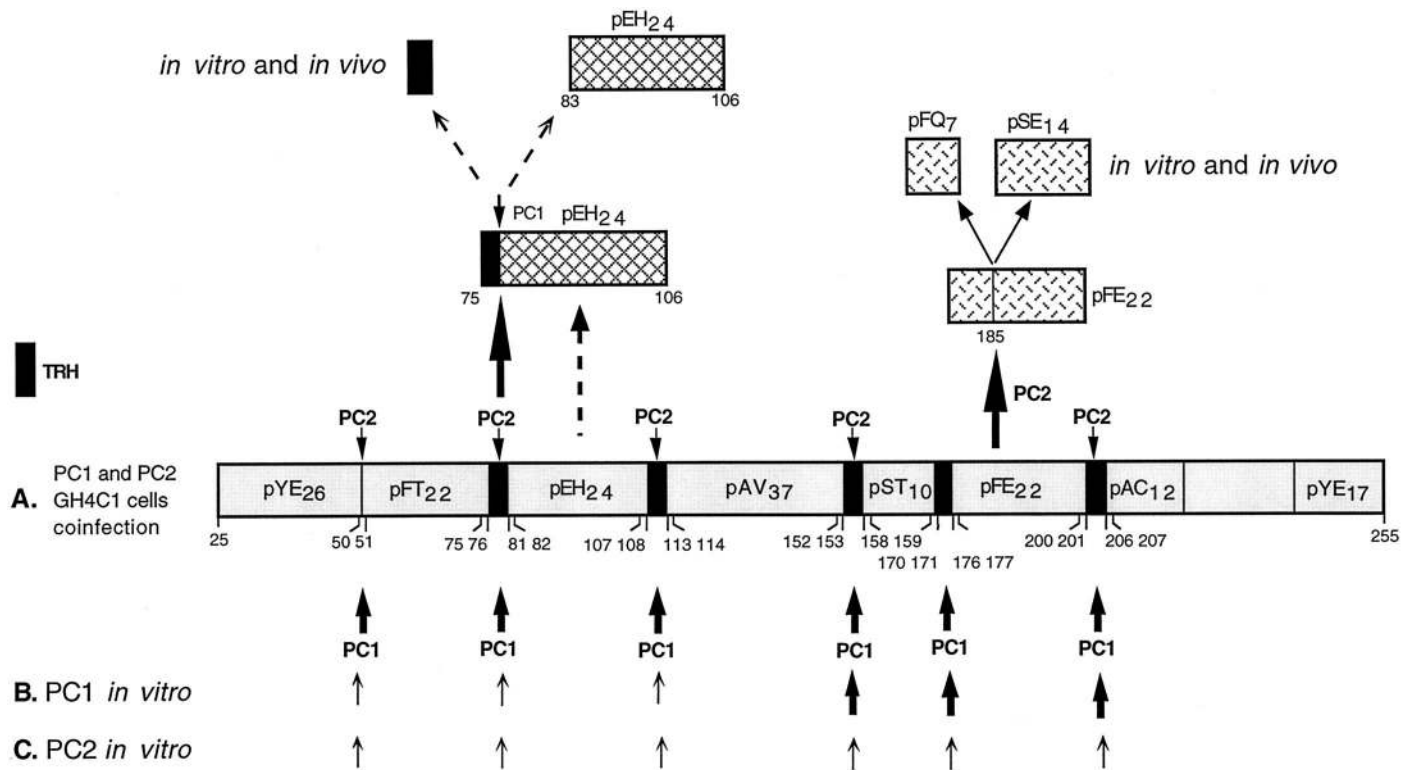


FIG. 6. Diagrammatic representation of rat pro-TRH and its cleavage by PC1 and PC2 as showed in our previous studies done *in vitro* and coexpression conditions (see Section II.D). The arrows indicate the site of cleavages and whether they are major (thick arrow) or minor (thin arrow) sites for each enzyme. [A portion of this figure was reproduced with permission from P. Schaner *et al.*: *J Biol Chem* 272:19958–19968, 1997 (37).]

products could be that TRH is extracted less efficiently from cells than these other peptides, although this seems unlikely because TRH is the smallest of the peptides, and doping experiments indicate TRH is extracted with high efficiency under our conditions. As described above, it is proposed that the processing of peptides derived from the N-terminal portion of pro-TRH is substantially different from the processing of those of the C-terminal end. This suggests the possibility of differential maturation of TRH molecules depending on location within the pro-TRH sequence (84). If this is the case, either the three TRH molecules derived from the C-terminal side, or the two from the N-terminal side, may play a more primary role in hormonogenesis. It is also possible that certain TRH molecules become biologically active at different times than others, or only some of them reach maturity while the rest are degraded by the TRH-degrading enzymes, depending upon physiological needs (see Section VII). Finally, cells may simply produce excess TRH that may or may not be used.

In summary, we have presented an overview of the current knowledge of pro-TRH biosynthesis, its processing, its tissue distribution, and the role of known processing enzymes in pro-TRH maturation. Evidence is presented suggesting differential processing for pro-TRH at the intracellular level is physiologically relevant. The data indicate that PC1 is primarily responsible for most pro-TRH cleavage events. PC2 is involved in specific processing events that occur later in the secretory pathway, specifically in the formation of the second TRH molecule from the N-terminal side of prepro-TRH_{83–106},

and the proteolytic cleavage of prepro-TRH_{178–199} to generate the novel prepro-TRH_{178–184} and prepro-TRH_{186–199} peptides.

E. Neuropeptide and catecholamine regulation of pro-TRH biosynthesis and processing

Immunoreactive TRH (iTRH) axon terminals are present in high density in the external layer of the rat ME, in close apposition to capillaries of the hypophysial-portal system (124). These axons originate from neuronal perikarya located in the PVN, the “thyrotrophic area” of the rat hypothalamus. Destruction of this region results in disappearance of up to 94% of TRH in the external layer of the ME and reduction of TSH secretion from the anterior pituitary gland (125). As described in Section IIC (Table 1), in addition to the PVN, iTRH neurons are present in other regions of the hypothalamus, including the POA, anterior hypothalamus, and supraoptic, arcuate, dorsomedial, and premmamillary nuclei, as well as basolateral and preformal hypothalamus (93).

Although many neurons in the PVN contain more than one peptide, TRH neurons are unique in being almost always unassociated with other known peptides (126). This makes the regulation of pro-TRH-derived peptide biosynthesis very specific. As described in more detail below, TRH neurons in the PVN are located in a region where they can be regulated by a number of neuroendocrine inputs. TRH neurons are densely innervated by norepinephrine (NE)-containing axons that stimulate TRH secretion (124). TRH neurons are also

densely innervated by neuropeptide Y (NPY) neurons. In smaller numbers, SRIF and endogenous opioid peptide (EOP) terminals are also in contact with TRH neurons (124). *In vivo*, these various neuroendocrine inputs may affect the levels of prepro-TRH mRNA and the posttranslational processing of PC1 and PC2. At the present time, the effect of this input on pro-TRH processing is unknown. Evidence for coordinated regulation of mRNAs for processing enzymes and their substrates has been documented in several cases (127). In contrast, outside the hypothalamus TRH is colocalized with other substances. For example, in the descending bulbospinal pathway, TRH is colocalized with SP and serotonin (5-HT) (128). Regulation of these TRH neuronal systems is much less well characterized than the thyrotropic neurons of the hypophyseal-portal system.

TRH-synthesizing neurons in the rat PVN receive a large number of afferent neuroendocrine inputs. Axon collaterals of parvocellular neurons ramify within the medial parvocellular PVN and establish numerous synaptic contacts with perikarya and dendrites of other parvocellular as well as magnocellular neurons, *e.g.*, SRIF afferents are derived from the PVN itself (129). The majority of inputs to TRH neurons are derived from the diencephalon, telencephalon, and brainstem (130). The paraventricular and medial parvocellular divisions of the PVN are densely innervated by NE-containing and epinephrine (E)-containing inputs from the medulla and pons (131). Further, NE-containing neurons densely innervate the midregion of the external layer of the ME. These inputs activate tuberoinfundibular neurons. Intracerebroventricular (icv) injections of NE, E, and α 2-adrenergic agonists stimulate basal TSH secretion (132, 133), and NE/E treatment of hypothalamic preparations stimulates TRH release (134). Inhibitors of catecholamine (CA) biosynthesis or α 2-adrenergic antagonists lead to a fall in basal TSH secretion. Thus, NE and E exert a tonic, stimulatory regulation on TSH secretion principally through α 2-adrenergic receptors. Stimulated release from the ME appears to be postsynaptically mediated via α 1-adrenergic receptors (135). In contrast, locus coeruleus (LC) afferents are inhibitory, being activated during stress (136). NE/E excitation of PVN TRH neurons mediates the rise in TSH in response to acute cold exposure or hypovolemia (135, 137, 138). However, it has also been proposed that α 1-adrenergic receptors mediate a phasic inhibitory regulation of TSH release. The data on NE/E modulation of TRH biosynthesis may be reconciled by an examination of how these inputs affect the posttranslational processing of pro-TRH, as well as examining their effects on PC biosynthesis. Peripheral levels of T_3 , T_4 , or TSH may also influence NE/E effects on TRH biosynthesis and/or release.

The PVN also receives prominent dopamine (DA) inputs from the posterior and dorsal areas of the hypothalamus, the zona incerta of the subthalamic region, and the A14 region of the anterior hypothalamus (124). Mesencephalic A9 and A10 dopaminergic neurons also project to the PVN. In turn, large terminal fields to the ME originate in the arcuate nucleus and periarculate nucleus regions of the hypothalamus. In contrast to the NE/E system, DA inputs appear to inhibit TRH secretion, mainly at the level of the ME (139). Aug-

mentation of DA neurotransmission inhibits basal and/or cold-stimulated TSH release, while DA antagonism has the opposite effect, although some studies have failed to replicate these findings (124). In addition, TRH release may be indirectly inhibited by DA-stimulated secretion of SRIF (140). Conversely, DA stimulates TRH release from isolated hypothalamic fragments (134, 140), again reinforcing the need to examine TRH biosynthesis in both *in vitro* and *in vivo* systems. Within the HPT axis, thyroid hormones appear to modulate DA levels in the ME, and TSH increases the ability of DA to inhibit TRH (141) (Table 1).

A wide array of neuropeptides, including NPY, TRH itself, SRIF, EOPs, neurotensin (NT), and vasoactive intestinal polypeptide (VIP), have inputs to the PVN and/or external layer of the ME (142–146). Other mediators, including γ -aminobutyric acid (GABA) and various cytokines, also appear to regulate TRH or TSH secretion, but there is as yet no anatomical evidence to support a direct action on TRH neurons in the PVN and/or ME (124). Anatomically, NPY appears most prominent in its inputs to the periventricular and medial parvocellular divisions of the PVN (147). NPY cell bodies principally reside in the medulla, often coexisting with NE and E (148), but other sources come from throughout the brain, including the arcuate nucleus of the hypothalamus itself. Indeed, the arcuate nucleus is the major source for NPY fibers innervating the TRH neurons in the PVN (149). Few NPY-containing axons project to the ME. The effects of NPY on tuberoinfundibular TRH are not yet well understood. NPY neurons also innervate SRIF neurons in the PVN, which would allow indirect regulation of TRH biosynthesis or secretion (124). Central administration of NPY reduces NE utilization in the PVN, as well as TSH release, indicating an inhibitory influence. *In vivo*, NPY also increases hypothalamic DA content, as well as DA turnover in the ME, the net result of which would reduce TRH release as well (150). Physiologically, NPY is critical to integrating thyroid function, food intake, and thermoregulation (151) (Table 1).

Inputs containing EOPs represent a second rich innervation to the PVN. These originate in the arcuate nucleus, periarculate area, and amygdala (144). The dorsal raphe projects 5-HT/enkephalin (ENK) axons, and the posterior hypothalamus-mammillary bodies send GABA/histamine/ENK projections, to the PVN. The ME contains numerous ENK, dynorphin, and endorphin (END) synapses originating from the PVN and arcuate nucleus. Both END and ENK inhibit TRH release from the hypothalamus, and ENK and morphine inhibit TRH secretion from the ME (152). There is additional evidence that ENK indirectly inhibits tuberoinfundibular TRH via DA release (153).

Recent data indicate that pro-TRH processing is regulated by opiate withdrawal (27). Opiate withdrawal increases prepro-TRH mRNA, and the N-terminal prepro-TRH_{53–74} and prepro-TRH_{83–106} peptides, in the rat PAG, whereas the level of TRH is unaltered (27, 154). New data also show suckling increases the production of prepro-TRH_{178–199} and prepro-TRH_{186–199} (see Section IVC and E). (91). These results demonstrate that levels of various products derived from pro-TRH can be posttranslationally regulated in an independent fashion under altered physiological conditions. Thus, it is

logical that neuroendocrine inputs into the PVN can affect pro-TRH processing as well.

Finally, we note that while the genomic organization of the rat prepro-TRH gene is well described (155, 156), the molecular mechanisms regulating the expression of this gene are incompletely understood. The 5'-region of the prepro-TRH gene contains TATA and GC box sequences, also present in the promoter region of other neuropeptide genes (155). In addition, sequences similar to a cAMP response element (CRE), and negative thyroid response elements (TREs), are present. The region between -47 and +6 of the rat prepro-TRH gene is active in CA77 TRH-secreting medullary thyroid carcinoma cells (155, 157), but not in transgenic mice (158). Inclusion of most of exon 1 (bp -47 to +84) increases promoter activity in CA77 cells and activates the promoter in transgenic mice, principally in prepro-TRH gene-producing tissues. Thus, *cis* element(s) located within exon 1 are necessary for the expression of the rat prepro-TRH gene *in vivo* (158). In CA77 cells, the human prepro-TRH gene is regulated by thyroid hormone through two distinct classes of negative TREs (157), similar to other neuropeptide genes such as prepro-SRIF (159).

F. Glucocorticoids modulate the biosynthesis and processing of pro-TRH

Glucocorticoids evoke a broad spectrum of responses in many eukaryotic cells by stimulating or repressing the transcription of glucocorticoid-regulated genes, including those of peptide hormones (160). The primary effect of glucocorticoids on gene transcription can occur by specific binding of the steroid receptor complex to DNA at the site of glucocorticoid response elements. Glucocorticoids can also interfere with the action of other transcription factors through protein-protein interactions and may elicit secondary effects at the posttranscriptional, translational, and posttranslational levels (161-163). For example, glucocorticoids stimulate pro-

cessing of the precursors for atrial natriuretic factor and neurotensin (NT) (162, 163). Glucocorticoids also regulate the posttranslational maturation, the intracellular trafficking, and the extracellular release of the mouse mammary tumor virus (164).

Glucocorticoids enhance TRH gene expression in several *in vitro* cell systems, including hypothalamic neurons, anterior pituitary cells, and thyroid C cells, an effect that occurs, at least in part, through transcriptional activation (165). Dexamethasone substantially elevates biosynthesis of the 26-kDa TRH prohormone and its intermediate products in cultured anterior pituitary cells, consistent with an overall up-regulation of both the biosynthesis and processing of the TRH precursor (161). This explains why glucocorticoids act not only at the transcriptional level, but also at the translational/post-translational level. This question can be addressed in experiments with AtT₂₀ cells transfected with prepro-TRH cDNA driven by a CMV-IE promoter not responsive to physiological signals. Dexamethasone causes a 75% increase in newly synthesized 26-kDa pro-TRH without altering prepro-TRH mRNA levels, suggesting that glucocorticoids raise translation rates and/or slow processing of pro-TRH. In fact, dexamethasone treatment accelerates TRH precursor processing.

Interestingly, processing of the N- vs. the C-terminal intermediates in the AtT₂₀ cells is influenced differentially by glucocorticoids. Levels of the N-terminally derived peptide prepro-TRH₂₅₋₅₀ are enhanced while levels of the 5.4-kDa C-terminally derived peptide are reduced. TRH content is increased (Fig. 7) (161). How could dexamethasone differentially affect the processing of N- vs. the C-terminal intermediates? Glucocorticoids may alter pro-TRH processing through changes in the expression of processing enzymes, as well as morphological alterations in AtT₂₀ cells. For example, GC volume is obviously enlarged in AtT₂₀ cells treated with dexamethasone. Although speculative, these changes may

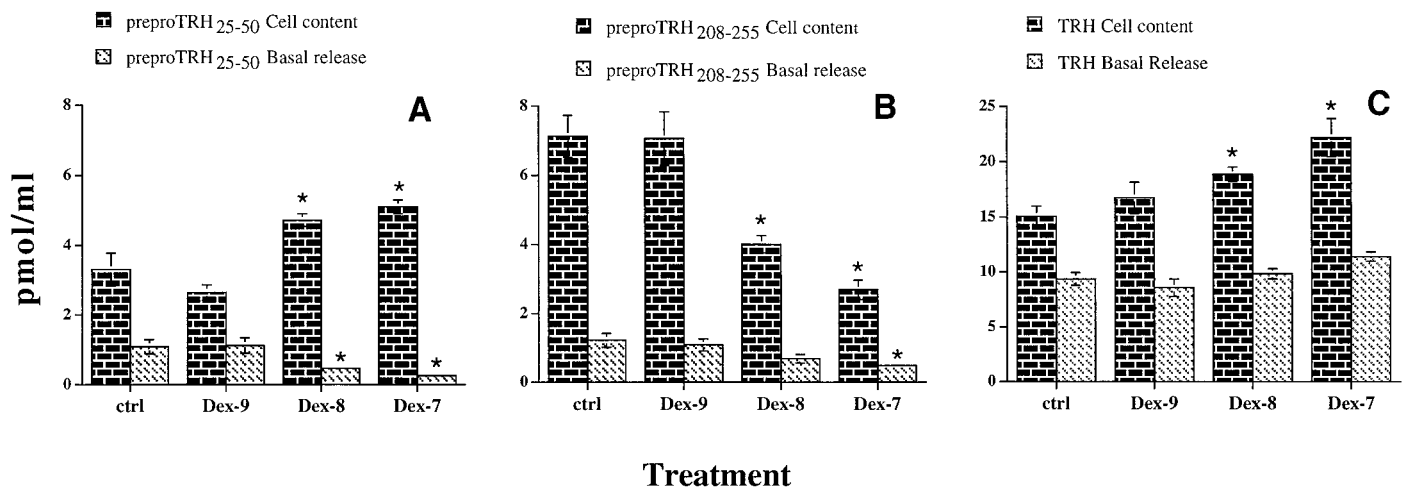


FIG. 7. Effect of dexamethasone on the accumulation and release of pro-TRH-derived peptides in AtT₂₀ cells transfected with a prepro-TRH cDNA. AtT₂₀ cells were cultured in six-well plates; cellular extracts and release media (2 h basal release) were processed further for the determination by RIA of the intracellular accumulation and basal release of the N-terminal peptide prepro-TRH₂₅₋₅₀ (A), the C-terminal peptide prepro-TRH₂₀₈₋₂₅₅ (B), and TRH itself (C) whose five copies are derived from both N- and C-terminal intermediates. A representative experiment (n = 6 wells for each group) is depicted. Data are presented as mean values ± SEM. *P < 0.05 compared with control. [Reproduced with permission from T. O. Bruhn *et al.*: *Endocrine* 9:143-152, 1998 (161).]

slow down the normal transport of intermediate products from the TGN to ISGs, thereby altering the accumulation or degradation of intermediate forms through changes in processing enzyme exposure (84, 161).

In vivo, more diverse effects of glucocorticoids on TRH gene expression have been reported. prepro-TRH mRNA-expressing neurons outside the PVN do not appear to be affected by changes in adrenal status, in contrast to the suppression caused by glucocorticoids in the PVN (166). The action of glucocorticoids on PVN prepro-TRH mRNA-expressing neurons may involve the hippocampus and amygdala, which convey negative feedback by glucocorticoids on CRF-expressing neurons in the PVN (167). Thus, direct positive regulation by glucocorticoids may be overridden by an indirect negative regulation causing a net reduction of prepro-TRH mRNA in the PVN while positive and negative regulation may sum to no change prepro-TRH mRNA-expressing neurons outside of the PVN (166).

In conclusion, glucocorticoids induce changes in the biosynthesis and processing of pro-TRH by affecting both transcription and translation rates, and by differentially influencing the processing of N- vs. C-terminal intermediates of pro-TRH. At the translational and posttranslational level, these effects result in an increase in TRH production, with more complicated differential effects on the accumulation of other N- and C-terminal pro-TRH-derived peptides. It is clear that control over the diverse range of pro-TRH-derived peptides within a specific cell is accomplished mostly from the regulation at the posttranslational level rather than the translational or transcriptional levels. Three examples supporting this hypothesis are presented in this review: 1) pro-TRH processing in the PAG is regulated during the opiate withdrawal, so that levels of TRH remain unchanged, but other pro-TRH-derived peptides are induced (*Section II.B.6*); 2) pro-TRH processing is regulated during suckling, where a selective, yet dramatic, increase in prepro-TRH₁₇₈₋₁₉₉ and prepro-TRH₁₈₆₋₁₉₉ peptides is observed (*Section IV.D and E*); and 3) in the absence of transcriptional effects, glucocorticoids induce differential processing of pro-TRH in both primary cultures of pituitary cells and transfected AtT₂₀ cells encoding prepro-TRH cDNA (this section).

G. Leptin regulates pro-TRH biosynthesis

Food deprivation in animals and humans results in endocrine and metabolic changes including decreases in circulating thyroid hormones, TSH, insulin, GH, gonadal hormones, and gonadotropins. Previous work in starved rats has shown a decrease in hypothalamic, but not thalamic, reticular, prepro-TRH mRNA, as well as decreased circulating TRH. This supports the concept that hypothyroidism produced after starvation is of hypothalamic origin (168). Leptin is a recently discovered peptide hormone that is synthesized and released by adipose tissue. Leptin also is decreased in starvation. Absence of leptin is responsible for the obese phenotype of *ob/ob* mice, and administration of this hormone to these animals decreases plasma corticosterone, suggesting that leptin is capable of inhibiting the hypothalamic-pituitary-adrenal axis. In normal rats and mice, leptin inhibits hypothalamic CRH release (169).

Leptin may have an important role in the neuroendocrine regulation of the HPT axis (170). During prolonged fasting in rats, low levels of T₃ and T₄ are observed, and TSH is in the low to normal range. As described above, this is due in part to suppression of prepro-TRH gene expression in PVN neurons. Since the decrease in thyroid hormone levels is blunted in mice and rats by systemic leptin, it has been proposed that the decrease in leptin detected during fasting alters the set point for feedback inhibition by thyroid hormones on the biosynthesis of prepro-TRH mRNA (170). The mechanism of such leptin regulation of prepro-TRH biosynthesis is unknown. It is hypothesized that leptin has direct actions on cell bodies in the arcuate nucleus, positively regulating POMC, and thus α -MSH, and negatively regulating NPY and the Agouti-related peptide (151). NPY afferents on TRH neurons are inhibitory (see *Section II.E*). In preliminary studies done in this laboratory, both leptin and α -MSH elevate prepro-TRH mRNA, pro-TRH, and TRH secretion in primary hypothalamic cultures (our unpublished results). Using the same primary cultures of hypothalamic neurons, leptin dose-dependently increases pro-TRH synthesis and TRH secretion. Immunocytochemical analysis reveals that approximately 40–50% of the hypothalamic cells are positive for the leptin receptor. Of these, approximately 10–15% colocalize with TRH (Fig. 8). These data suggest that the regulation of pro-TRH biosynthesis and TRH release in response to starvation includes direct regulatory actions of leptin and α -MSH on hypothalamic TRH neurons involved in HPT axis homeostasis (171).

In summary, leptin effects on pro-TRH biosynthesis include: 1) an inhibitory action of leptin on NPY release from the arcuate nucleus, which in turn may reduce the inhibitory action of this peptide on TRH release from the PVN; 2) a stimulatory action of leptin on α -MSH release from the arcuate nucleus, which may stimulate TRH release from the PVN; and 3) a direct action of leptin on TRH neurons located in the PVN.

III. Function of TRH

A. The HPT axis

The hypophysiotropic role of TRH in the control of thyroid function has been extensively reviewed elsewhere, and the reader is referred to the excellent review by Scanlon and Toft (172). Therefore, this section will only briefly describe the role of TRH in the HPT axis, including relevant new reports, to aid in the understanding of other sections of this review.

TSH is synthesized and secreted by the thyrotrophic cells in the anterior pituitary and is the major regulator of the thyroid gland. TSH secretion is primarily regulated by negative feedback from circulating thyroid hormone and by stimulatory input from the hypothalamus through TRH action on thyrotrophs. There are other factors known to regulate TSH secretion, including glucocorticoids in the systemic circulation and SRIF and DA from the hypothalamus. There is evidence supporting the view that glucocorticoids in man suppress endogenous hypothalamic TRH secretion (173). Further support for this hypothesis was demonstrated in adrenalectomized rats in which pro-TRH mRNA levels

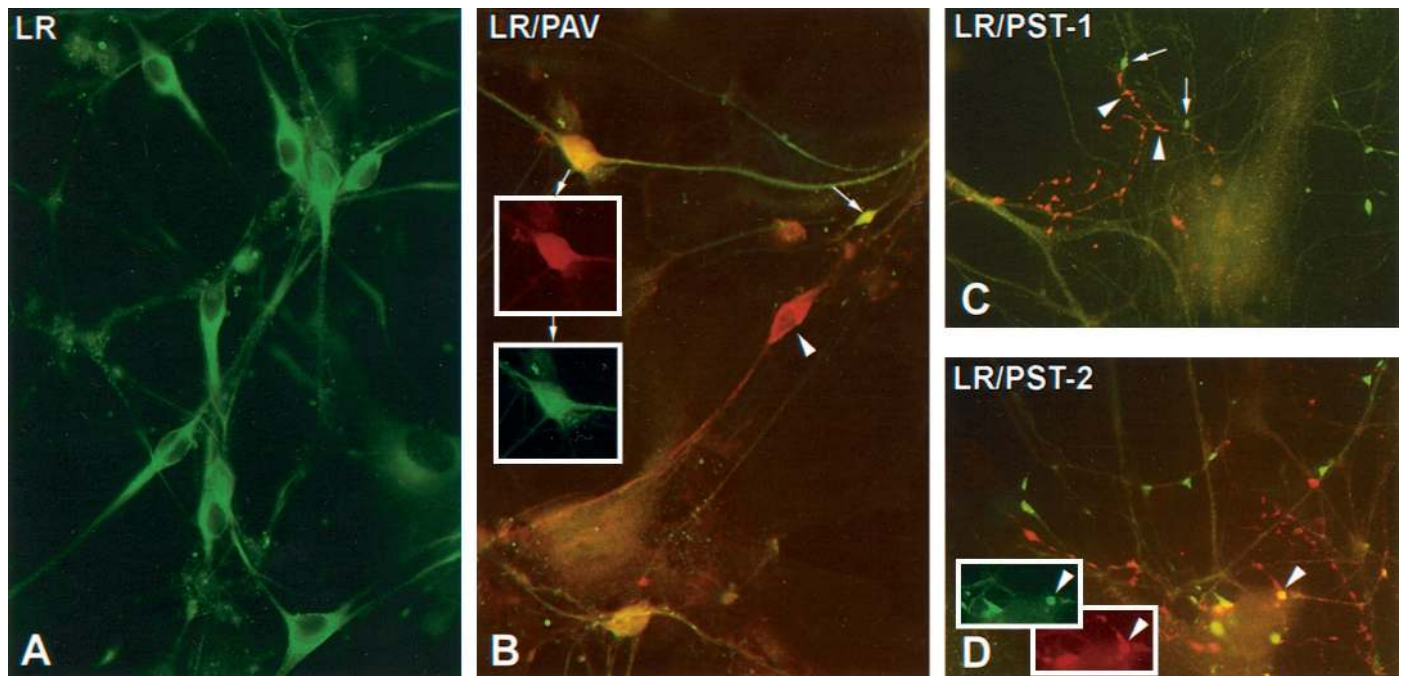


FIG. 8. Colocalization of pro-TRH with the leptin receptor (LR). Neuronal cells cultured for up to 14 days in four chamber labTeck slides were fixed with 4% paraformaldehyde followed by immunoreaction with anti-LR and anti-pro-TRH. Fluorescein isothiocyanate conjugated to goat antirabbit globulin was used as a probe. Texas Red-X-succinimidylester directly conjugated to anti-pAV₃₇ or anti-pST₁₀ antibodies were used as a probe for pro-TRH. Panel A shows positive immunostaining for LR in neurons untreated with Triton X-100. Staining is mainly localized on the surface of cell body and dendrites. Panel B shows cells with positive immunostaining for pro-TRH using anti-pAV₃₇ (red color, arrowhead) and LR (green color, arrow). Colocalization of pro-TRH with LR (yellow-orange color indicated). The insets show the same cell with each individual staining. A distinct population of neuronal cells contains pro-TRH but not LR and vice versa. Among the total cells with positive staining for pro-TRH, approximately 30% or less of those cells colocalize with LR. Panel C shows positive immunostaining for pro-TRH using anti-pST₁₀ (arrowheads, red) and for LR (arrows, green), which are present in different subcellular locations. Panel D shows positive immunostaining for pro-TRH peptides (red color, arrowheads) that colocalize with LR (green color). Thirty millimeter slides were digitized with a video camera and appropriate macro lens using BioVisionframe grabber software (Perceptics Corp., Knoxville, TN). Images of the red and green planes were combined using Adobe Systems to show areas of colocalization. The resulting images were printed with a Mitsubishi CP210 dye sublimation printer (Apunix Computer Services, San Diego, CA).

increase 1.6-fold, an increase reversible with dexamethasone (166). The role of glucocorticoids *in vivo* and *in vitro* has been described in the previous sections. Both SRIF and DA inhibit TSH release by direct effects on thyrotrophs. As discussed in Section II.C., SRIF perikarya in the POA, periventricular parts of the anterior hypothalamus, and a few in the PVN, and TRH neurons in the PVN, project to the ME. TRH and SRIF are the two main TSH-regulating hypophysiotropic neurohormones released into the hypophysial portal vasculature. The maintenance of euthyroidism is dependent on a highly regulated balance of neuropeptides and neurotransmitters, where the dominant positive hypothalamic control for TSH is TRH, and the principal feedback control is through thyroid hormones. However, thyroid hormones also provide direct negative feedback on prepro-TRH gene expression as well. The relationship of thyroid hormone regulation to pro-TRH processing is undefined.

Even though TRH is the major regulator of the synthesis and secretion of TSH, and thus plays a pivotal role in the HPT axis, in a recent study, homozygous TRH gene knockout mice were shown to be viable, fertile, and exhibit normal development (174). Whereas the TRH^{-/-} mice showed normal serum PRL and GH levels, thyroid hormone levels were significantly reduced as compared with the wild-type heterozygous mice. The targeted disruption of the prepro-TRH

gene caused a characteristic tertiary hypothyroidism, and a substantial decrease in insulin secretion resulting in a profound hyperglycemia. These authors suggested that in addition to abnormalities of the thyroid function, TRH may be involved in the pathogenesis of diabetes mellitus (174).

B. Extrahypophysiotropic TRH

More than two-thirds of iTRH in the brain is found outside of the traditional "thyrotrophic zone" of the hypothalamus (175, 176). This extrahypophysiotropic TRH is believed to function as a neuromodulator of known neurotransmitters (177, 178). Indeed, it might act as a neurotransmitter itself; it is present in secretory granules whose exocytosis is responsive to membrane depolarization, it acts through specific receptors that are widely distributed throughout the CNS, and it is rapidly cleared through specific catabolic pathways (179).

While the following discussion focuses on TRH, many other neuropeptides and neurotransmitters play critical roles in the biological functions discussed below. In several areas of the brain, TRH is colocalized with other neurotransmitters and/or neuromodulators, including 5-HT and SP in the bulbospinal pathway, DA in the olfactory bulb, and histamine, ENK, and NPY in various loci of the hypothalamus (128).

Where TRH is directly affected by, or directly affects, other neuroactive agents, they have been discussed, but otherwise their roles are left for review elsewhere. Further, in the following we describe many effects of TRH. In fact, these have been demonstrated using TRH and/or TRH analogs. Due to space constraints, we will not distinguish effects by TRH *vs.* its analogs. Available TRH analogs have higher affinities for the TRH receptor, longer half-lives, etc. and are reviewed elsewhere (180, 181).

Finally, for several TRH effects, the metabolite histidylproline diketopiperazine, or cyclo (His-Pro) (CHP) also has agonist or antagonist effects. CHP is present in the CNS and peripheral tissues in levels that often exceed levels of TRH (182). While CHP is a known breakdown product of TRH, there are data that CHP is also derived from precursors other than TRH (183). Further, high-affinity binding sites for CHP have not been identified (184) (but see Ref. 185). Thus, the precise biological meaning of CHP effects is unknown. Figure 9 summarizes proposed physiological roles for extrahypophysiotropic TRH.

1. TRH and ergotrophic effects. In diverse ways TRH is excitatory when infused into animals (179). Activation is reflected in organism-wide effects on arousal, sleep, cognition, locomotion, and mood (186). Metcalf and Dettmar (187) first used Hess's term "ergotrophic" to describe the endogenous activating effects of TRH in the brain that are described below.

a. TRH in arousal and sleep: Systemic (188, 189) and central (190) TRH increase wake time and/or decrease sleep time in multiple species. More dramatic is the ability of TRH to arouse animals from drug narcosis induced by alcohol (191, 192), β -endorphin (193), tetrahydrocannabinol (THC) (194), benzodiazepines (195), and barbituates (196, 197). icv Anti-TRH antiserum doubles anesthetic-induced sleep time, supporting an endogenous role for TRH in arousal (198). Of unclear significance, CHP is more effective than TRH in

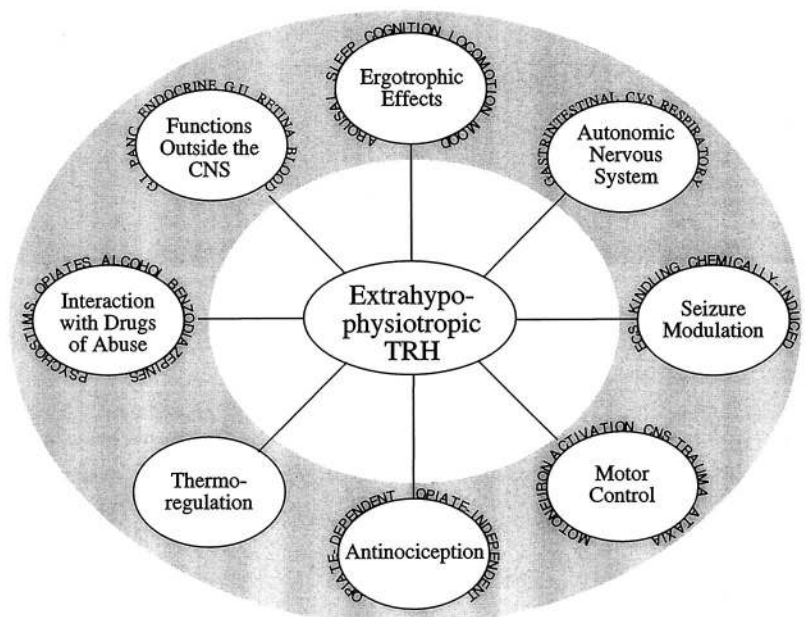
decreasing sleep time and reversing ethanol-induced narcosis, but does not affect TRH modulation of barbiturate-induced sleep time in mice (199, 200).

TRH analepsis is most strongly induced by infusions into the medial septal area, the diagonal band of Broca, or the nucleus basalis of Meynert (186, 201). TRH levels rise in the medial septum as rats recover from ethanol-induced sedation (202). Further, TRH enhances cholinergic activity in the septo-hippocampal and nucleus basalis-cortical systems (203–205), pathways that play a central role in reversal of drug-induced narcosis (206). Atropine blocks both analepsis and cholinergic enhancement when TRH is infused into these areas. However, antagonism of the analeptic response to *systemic* TRH requires both ACh and NE blockade, supporting mediation by other neuroanatomical sites (192, 207). The posterior lateral hypothalamus, the midbrain reticular formation, and TRH-containing fibers passing through, or synapsing in, the dorsal septum are additional neuroanatomical substrates that might mediate arousal through noradrenergic mechanisms (203).

There are limited data that TRH may be useful to enhance arousal in pathological states. TRH prevents the postconcussive elevation of cortical ACh and reduction of cortical NE seen in mice (208) and decreases time of unconsciousness in head-injured mice (209). Canine models of narcolepsy show improvement after treatment with TRH (189).

b. TRH in cognition: The ergotrophic effects of TRH on consciousness and arousal are often detected along with improvements in measures of learning and memory, consistent with the important role of ACh in both processes (210). TRH improves performance in learning-impaired mice, an effect that is blocked by scopolamine (211). TRH restores learning and/or memory deficits in rats made cognitively deficient by anticholinergic treatment, electrochemical shock treatment (ECS), or surgical lesions (212–214). These effects are largely mediated by enhanced cortical ACh

FIG. 9. Physiological roles of extrahypophysiotropic TRH. Extrahypophysiotropic TRH is implicated in a wide range of physiological processes, as well as processes affected by drugs of abuse. These functions are broken down into areas portrayed as ellipses within the darker annulus surrounding extrahypophysiotropic TRH in the center. Major subjects discussed in the text for each area are shown *outside the ellipses*.



release with TRH infusion into the nucleus basalis of Meynert and increased hippocampal ACh with TRH infusion into the medial septum-diagonal band (215). Facilitated release of NE (216) and *N*-methyl-D-aspartate (NMDA) receptor activation (217) also are implicated in cognitive enhancement by TRH. Human trials have demonstrated only modest cognitive improvements in patients with alcoholic dementia (218), in the ECS postictal state (219), and in Alzheimer's disease (220, 221).

c. TRH in locomotor activation: Systemic TRH elicits a number of motoric and behavioral activating effects in a wide range of species (178, 222). Further, TRH antagonizes locomotor depression induced by alcohol (223) and β -endorphin (193). TRH elicits locomotor activation when injected into the NAc (224, 225), the ventral tegmental area (VTA), the caudate (222), the septal nuclei (225), and the ventromedial hypothalamus (226).

Locomotor activation by TRH principally is mediated by DA. Repeated TRH treatment in rats elevates DA in the cerebral cortex and increases tyrosine hydroxylase activity, the rate-limiting enzyme in DA biosynthesis in the brain stem. These effects correlate with dose- and time-dependent increases in locomotor activity (227). DA antagonists and/or DA depletion block locomotor activation by systemic TRH (228). Consistent with TRH activity on the mesocorticolimbic system, systemic TRH increases DA release in the NAc (224, 229), and bath application of TRH induces DA release from NAc (230) and septal (231) brain slices.

Intraaccumbens TRH increases DA metabolism in the NAc, while icv TRH or intra-VTA TRH do not, although these latter treatments elicit locomotor excitation (222). Further, TRH can stimulate locomotion at doses that have no effect on NAc DA (232). These data support redundant but distinct TRH mechanisms, in addition to those in the mesocortical DA system, that induce locomotion (233). Other studies indicate that opiate antagonists and α -adrenergic antagonists can attenuate TRH-induced locomotion (234). A single study reports that CHP is able to stimulate locomotion (200).

d. TRH and antidepressant effects: In a number of behavioral assays used to screen compounds for antidepressant efficacy, TRH tests positively (235). This potentiation is independent of effects on TSH or thyroid hormones; however, PRL similarly can potentiate desipramine effects in the forced swim test (236). For the most part, DA receptor antagonists block the antidepressant effects of TRH (237), although it is difficult to tease out locomotor effects from antidepressant effects in some behavioral tests. TRH action in some antidepressant screens is blocked by opioid receptor antagonists or α 2-adrenergic blockade (237). Further, antidepressant treatments alter TRH levels in rat brain, but not in a clear "antidepressant pattern" (238). In human trials, promising early results (239, 240) have given way to larger studies that indicate TRH is of limited benefit in depression (241, 242). Recent studies using intrathecal TRH reveals significant reductions in symptomatology of patients with refractory depression, although tachyphylaxis to the effects develops rapidly (243, 244). Recently, a compelling model has been put forward that clinical depression results from pathologically

overdriven glutamatergic circuits in the limbic forebrain that have escaped inhibitory regulation by TRH (245).

2. TRH and autonomic nervous system function. The brainstem distribution of TRH supports a prominent role in autonomic nervous system (ANS) function. Fully 65% of medullary TRH is associated with dorsal vagal complex (DVC) neurons of the nucleus tractus solitarius (NTS), nucleus intercalatus and commissuralis, the dorsal motor nucleus (DMN) of the vagus, and, to a lesser extent, the nucleus ambiguus (246). Injection of TRH onto DMN neurons is uniformly excitatory, while applications onto NTS neurons are inhibitory (247). The majority of DVC iTRH derives from fibers arising from the medullary raphe nuclei that pass through the DVC (93, 248). However, cells within the DMN express prepro-TRH mRNA and pro-TRH (95, 249), consistent with some endogenous TRH production.

A descending bulbar-spinal pathway, in particular, from the nucleus interfascicularis hypoglossi and the nucleus paragigantocellularis lateralis, projects to the intermediolateral (IML) column of the spinal cord (250). Fibers and terminals of this tract are closely apposed to preganglionic sympathetic neurons. TRH fibers and preganglionic sympathetic neurons are also found around the central canal and in the intermediate gray matter of lamina VII (93, 251). While some studies indicate that more than 90% of TRH immunoreactive neurons also stain positively for 5-HT, and 75% express immunoreactive SP (252), only 43% of IML TRH is ablated by 5-HT neurotoxins (250). Indeed, more than 90% depletion of 5-HT in the spinal cord reduces spinal cord TRH by only 66% (253). Thus, a sizable pool of TRH-containing neurons are not serotonergic.

a. TRH and gastrointestinal function. TRH inhibits food and water intake: TRH inhibits food and water intake, consistent with its high levels in the ventromedial hypothalamus (176, 254), a center important to regulation of food intake (255), and its interaction with NPY and NE, both important to intake behavior (93). Systemic TRH reduces food intake less effectively than icv TRH, arguing for a central effect (256). Parenteral TRH can suppress eating without altering blood glucose levels (257) and without affecting TSH (256). icv TRH also reduces water intake (258), although others have reported icv TRH reduces food intake far more than water intake (259). icv TRH suppression of stress-induced eating is antagonized by D-alanine-enkephalin (260), although DA transmission in the nigrostriatal pathway and lateral hypothalamus also affects stress-induced eating (261).

The hypothalamus serves as a principal brain substrate to coordinate hunger and satiety; it is generally held that the ventromedial hypothalamus serves to signal satiety, and the lateral hypothalamus, hunger (255). Injection of TRH into the ventromedial hypothalamus is most potent in producing adipsia and anorexia, and lateral hypothalamus injection is selectively potent for adipsia (262). Iontophoretic application of TRH onto ventromedial hypothalamic neurons results in facilitation of glucoreceptors, and hence, decreased feeding drive (263). However, others argue that the lateral, not ventromedial, hypothalamus is most critical for TRH-induced anorexia (226). TRH administration into the NAc is mildly anorexic (262). Systemic and icv CHP also suppress sponta-

neous food and water intake and stress-induced feeding, although it is less potent than TRH in feeding (264). In water-deprived rats, CHP is equipotent with TRH in reducing drinking (265). Thus, it cannot be ruled out that the TRH effects discussed above are actually the result of CHP as a TRH metabolite.

TRH enhances gastric acid secretion: Vagal preganglionic neurons arising from the rat DVC and nucleus ambiguus terminate in the gastrointestinal tract (266, 267), in close proximity to nerve efferents of the greater curvature and pylorus of the stomach (268). These comprise the major medullary projections to the stomach. In the cat, retrograde tracing does not support a descending tract from the nucleus ambiguus, but the remainder of the DVC participates in the bulbogastric projection (268). The high concentration of TRH receptors in the DVC (269, 270) in close proximity to NTS vagal afferents (266) is consistent with modulation by TRH afferents from the NTS and the medullary raphe nuclei. As well, peripheral signals of gastric distention and gastric secretion are carried by vagal afferents back to the medulla to activate DVC neurons. The codistribution of afferent and efferent pathways in the vagus provide a means for bulbogastric TRH neurons to modulate gastrointestinal responses to physiological signals, such as gastric distension, the cephalic phase of gastric acid secretion, etc. (271, 272).

Central TRH is far more potent than iv TRH in inducing gastric acid secretion (18, 273). icv Anti-TRH inhibits gastric acid secretion in pylorus-ligated (274) and cold-restrained (18) rats, supporting an endogenous role for the peptide. TRH stimulates gastric acid secretion independent of hypophysiotropic effects or effects on gastrin (275, 276). Further, TRH injection into the DVC is 10 times more potent than icv TRH in stimulating gastric acid secretion (18, 277, 278). Bilateral DVC injection of anti-TRH antiserum significantly reduces gastric acid secretion in response to icv TRH (279), or chemical or electrical activation of medulla raphe pallidus (RPa) neurons (280), supporting the central role of the DVC in TRH effects on gastric acid secretion.

Atropine injection into the DMN does not completely block TRH-stimulated gastric acid secretion (279), because other loci, including the nucleus ambiguus, lateral hypothalamus, and the ventromedial hypothalamus, can mediate TRH-induced gastric acid secretion (18, 281). TRH action also is partly mediated through α_2 -adrenergic receptors and enhanced sympathetic outflow that modulates the vagus (275, 277, 282). By unknown mechanisms a number of peptides in the DVN, including CRF, bombesin, calcitonin gene-related peptide (CGRP), calcitonin (CT), endogenous opiates, and, curiously, gastrin releasing peptide, inhibit TRH-induced gastric acid secretion (18). CHP and TRH-OH have no such activity (277, 283).

Kainic acid stimulation of afferent nucleus raphe obscurus (ROb) neurons mimics the induction of gastric acid secretion by TRH injection into the DVC (278). In addition, the caudal raphe nuclei-DVC pathway mediates cold-induced vagal stimulation of gastric acid secretion and erosion formation (284). Further, anti-TRH antisera injected into the DMN abolishes the ability of excitatory amino acid injection into the RPa to enhance indomethacin-induced gastric erosion for-

mation (285). Thus, excitation of the raphe nuclei enhances DVC outflow, and one of the mediators of this effect is TRH. 5-HT (286) and SP (287) afferents from the raphe nuclei to the DVC modulate TRH effects.

icv TRH administration indirectly affects gastric acid secretion by increasing pepsin secretion and gastric mucosal blood flow and secretion (18). This effect is partly inhibited by DVC injection of anti-TRH antiserum, and surprisingly, is independent of increased gastric acid secretion (279). Thus, TRH may provide a means to regulate pepsinogen secretion without altering acid production. Intracerebral (ic) TRH-stimulated gastric mucosal blood flow is vagally mediated, via stimulation of an L-arginine-nitric oxide (NO) pathway independent of histamine H1 receptors or capsaicin-sensitive afferents (288). In addition, ic TRH enhances gastric secretion of 5-HT and 5-HT entry into the portal vasculature, an effect that again is vagally mediated (289).

TRH effects on gastrointestinal contractility and transit: ic, But not iv, TRH increases gastric contractions and gastric emptying in most species (290–292). Enhanced gastric motility is reproduced by direct infusion into the DMN but not the nucleus ambiguus (268), an effect completely blocked by vagotomy (293). Systemic morphine (294), or DVC injections of CRF (295), bombesin (296), and interleukin-1 β (IL-1 β) (297) inhibit the TRH effect. Since gastric contractility is inhibited by excitatory amino acid injection into the DVC (298), it is likely that TRH is inhibitory to DVC neurons controlling gastric contractility. Indeed, TRH injections onto NTS neurons reduce their spontaneous activity (299).

The ROb, RPa, and nucleus raphe magnus (RMg) provide afferents to the DVC (300, 301). In the cat, the DMN receives its strongest inputs from the caudal RPa and ROb, where TRH neurons are enriched (301). Glutamate or electrical excitation of the caudal RPa and ROb, but not rostral RPa or RMg, results in enhanced gastric contraction. This effect is abolished by vagotomy and anti-TRH antibody injection into the DVC (302). Surprisingly, TRH stimulates ROb and RPa TRH afferents to the DVC. This effect is completely abolished by vagotomy or atropine into the ROb, markedly attenuated by atropine into the RPa (303), and antagonized by SP or VIP into the ROb (304). ic Antisense oligonucleotides to the TRH receptor block the increase in gastric motility seen with TRH injection into either the ROb or the downstream DVC, while glutamate excitation is unaffected. Thus, TRH activates both TRH and cholinergic afferents to the DVC, which in turn increase gastric motility (305). Finally, gastric contractility also is increased by TRH injection into the hypothalamic paraventricular nucleus or the central nucleus of the amygdala, an effect abolished by subdiaphragmatic vagotomy. However, unlike medullary effects, the frequency of gastric contractions after these injections is attenuated (306, 307).

Motility in the proximal small intestine and ascending colon and cecum is also mediated by a central effect on vagal outflow (291, 308). However, central depletion of brain catecholamines blocks the contractile response in the duodenum, indicating a critical role for catecholamines as well as ACh in TRH central regulation of bowel motility, at least in some regions of the gut (309). Acceleration of small intestinal transit appears to occur through a separate pathway from

that described for the stomach (310). TRH (≥ 100 ng) increases small intestine transit only when injected into the medial septum, or lateral and anterior hypothalamus, in anesthetized rats. icv, But not iv TRH, also reverses net water absorption in the jejunum and ileum, an effect entirely abolished by vagotomy (311). Large colon transit in rabbits is increased by iv or icv TRH (312) and is associated with accumulations of fluid in the colon (289). TRH effects on colonic transit are mediated by vagal and sacral cord parasympathetic outflow, as well as serotonergic transmission (18). In humans, iv TRH retards glucose and xylose absorption by the gut (313).

TRH effects on pancreas and liver. In normoglycemic rats, acute systemic TRH will induce hypoglycemia, with little effect on peripheral pancreatic hormones (314). Central TRH potently blocks epinephrine-induced hyperglycemia, presumably via combined parasympathetic/sympathetic induction of insulin secretion (315). Central TRH antagonizes hyperglycemia induced by treatments other than epinephrine, including central injections of CRF, ENK, and glucagon, as well as systemic 2-deoxyglucose, foot shock, immobilization, or endotoxin (315).

Pancreatic effects of TRH are most likely paracrine. TRH is synthesized in the insulin-producing β -cells (316). In neonatal pancreas, TRH and insulin appear to be secreted via the same potassium-, cAMP-, and protein kinase C-responsive pathways (317); in adult pancreas, TRH secretion is inversely related to insulin secretion (318). While TRH does not affect insulin release (319), TRH and CHP inhibit 2-deoxyglucose-stimulated pancreatic secretion in a dose-dependent manner (320), and TRH enhances arginine-stimulated glucagon release (319). In isolated rat pancreas perfusates, anti-TRH antiserum reduces glucose- and arginine-stimulated glucagon secretion, and exogenous TRH enhances basal glucagon secretion if endogenous TRH is first cleared (318).

TRH also mediates central effects on pancreatic secretion. CSF injection of TRH, and microinjection into the DVC, stimulates exocrine pancreatic volume, protein, and bicarbonate secretion via vagal outflow (321, 322). VIP is also an important nonmuscarinic mediator of TRH-stimulated pancreatic secretion, while CGRP, via noradrenergic mechanisms, opposes TRH pancreatic stimulation (321). Curiously, in isolated rat pancreatic acinar cells, TRH inhibits carbachol- and ceruletide-stimulated, but not OAG- or CCK-stimulated amylase secretion (320, 323). Thus, certain pancreatic secretory pathways may show opposite peripheral and central effects by TRH. Finally, TRH has a trophic effect on the pancreas (324). Chronic administration of TRH for 10 days via gastric fistula significantly increases pancreatic weight, DNA content, and protein content, although enzyme concentrations are not proportionally elevated, so that their final concentrations are reduced.

ic RX77368, a TRH analog, stimulates hepatic DNA synthesis 24–72 h post injection in a dose-dependent manner (325). iv Administration is ineffective. The effect is abolished by hepatic vagotomy or atropine. Further, ic RX77368 enhances hepatic blood flow 15–90 min post injection. This regulation is abolished by hepatic vagotomy, atropine, in-

domethacin, or the NO synthesis inhibitor, NG-nitro-L-arginine methyl ester (326).

b. TRH and cardiovascular function: TRH reverses shock of varying etiologies in a number of animal species (327). However, the precise cardiovascular changes induced by TRH vary markedly with dosage, species, and experimental state, in particular, whether the animal is anesthetized or conscious, and normotensive *vs.* hypotensive. While TRH displays many cardiovascular effects, CHP and TRH-OH have minimal cardiovascular activity (328).

In rabbits, both anesthetized and conscious, iv TRH increases blood pressure and causes peripheral vasoconstriction (328, 329). Effects in anesthetized rats are the same, except peripheral vasodilation is seen (330); this peripheral vasodilation is reversed by cholinergic blockade (328). Overall, TRH modulates blood pressure through combined parasympathetic and sympathetic effects (328, 330, 331). DA plays a lesser role (332), and naloxone is ineffective in altering these effects (333). As well, iv TRH in rats completely reverses systemic NT-induced hypotension and attenuates the central pressor effect of NT (334).

In anesthetized rabbits, but not conscious rabbits, cerebral vasodilation is induced by iv TRH, bringing cerebral blood flow back to the level observed in conscious animals. Once cerebral blood flow is normalized, TRH has little effect (329). Cerebral vasodilation is not the result of alterations in peripheral blood pressure, blood gases, loss of autoregulation, change in cerebral metabolism, or change in oxygen saturation gradients (328). Cerebral vasodilation in anesthetized rabbits is partially blocked by the α 2-adrenergic antagonist yohimbine, although yohimbine has no effect on TRH-induced elevation of mean arterial pressure (335). Neither vagotomy nor cholinergic blockade reduces these effects (328). Transection at the mesencephalic pons abolishes TRH-induced cerebral vasodilation without affecting its systemic pressor effect (336). Cordotomy at the C1 level abolishes the pressor effect of TRH, but not its effect on the cerebral vasculature. Thus, peripheral pressor and vasoconstriction effects are mediated more caudally than those increasing cerebral blood flow.

icv TRH significantly elevates blood pressure and heart rate in anesthetized (337, 338) and conscious (339) rats. In anesthetized rats, icv TRH increases blood flow to most organs, and this is abolished by bilateral vagotomy (328). In conscious rats, much more peripheral vasoconstriction is found (340), although this effect is reversed in hypovolemic states (341). icv TRH elevates plasma levels of NE and E independently of plasma renin activity or vasopressin. The vascular effects of icv TRH in rats are mimicked by TRH activation of sympathetic nuclei within the brain (342); intrathecal TRH induces its pressor response in rats and humans via increased sympathetic activity to the peripheral vasculature and adrenals (343), an effect mediated in rats by TRH receptors in the IML (344). Interestingly, sympathetic nerve responses in rats are partially attenuated by reduced thyroid activity, providing a second route for TRH regulation of cardiovascular function (345). Taken together, the data in rabbits and rats suggest that central TRH regulation of cardiac functions and organ blood flow distribution principally

are mediated through the sympathetic nerves and the adrenal medulla (339).

Selective electrolytic lesions have identified the dorsal raphe nuclei as mediating the pressor response induced by icv TRH. Reductions in blood pressure induce prepro-TRH mRNA in the RPa (346). These changes, in turn, effect the descending bulbospinal tracts and ascending tracts to the PVN (342). TRH infusion directly into the dorsal raphe nuclei reproduces the pressor, tachycardic, and sympatho-excitatory effects of icv TRH, and these effects are blocked by ganglioplegia with pentolinium. Further, chemical lesioning by the 5-HT-preferring toxin 5,7-dihydroxytryptamine (5,7-DHT) obliterates TRH-induced tachycardia, while the NE/DA-selective toxin 6-hydroxydopamine does not. These data strongly implicate the descending bulbospinal pathway to the IML column to boutons on preganglionic sympathetic neurons as the neuroanatomical pathway mediating the peripheral cardiovascular effects of TRH (347).

TRH in experimental CVS disease: In experimental models of anaphylactic shock, hemodynamic parameters in mice (348) and guinea pigs (349) are improved by iv TRH, largely through elevation of plasma E and NE (350). icv TRH mimics the protective effect of iv TRH (351). TRH action in endotoxic shock also is mediated through the sympathetic nervous system (350). In hemorrhagic shock, TRH elevates blood pressure either by improvement in cardiac output, or by increasing peripheral vascular resistance (352). However, after effects of the latter may be harmful and could explain why TRH efficacy for survival after hemorrhagic shock has varied among species (353, 354). Although principally mediated by catecholamines, icv TRH-induced hemodynamic improvements in shock are partially blocked by δ -opiate receptor antagonism (354) and vagotomy or atropine sulfate (355).

Few studies have examined the converse effect of shock on TRH. In rats, hemorrhagic shock decreases TRH in frontal cortex, septum, hippocampus, and hindbrain (356). TRH receptor binding is significantly decreased in septum and hindbrain. However, other experiments indicate that elevations of TRH in the medulla, midbrain, cortex, striatum, and cerebellum after hemorrhage are associated with better survival (357). Thus it appears that reductions in TRH neurotransmission in certain parts of the brain may contribute to the pathophysiology of shock. Further, TRH effects may be mediated, in part, by downstream effects on the thyroid axis (358) and plasma vasopressin (204).

In spontaneously hypertensive rats (SHR), CSF TRH, prepro-TRH mRNA, TRH, and TRH receptor binding in the POA are all significantly elevated compared with Wistar-Kyoto (WKY) control rats (359). In SHR, elevated TRH receptor binding in striatum and hypothalamus correlates with the development of hypertension (360). Further, iv or icv TRH antiserum lowers arterial blood pressure in SHR rats, but not WKY rats. Chronic enalapril, a vasodilator, decreases blood pressure and reduces POA TRH levels, although another vasodilator, diltiazem, has no effect (359).

Animal models of stroke have provided a testbed for potential therapeutic application of TRH. In rats with middle cerebral artery (MCA) occlusion-induced infarcts, icv TRH given at 15 min and 24 h post surgery significantly improves

survival, protects against ischemic damage, and reduces infarct size 10 days after surgery (361). Also in MCA infarcts, TRH increases blood flow to the infarct area (362). ip And oral TRH improve neurological deficits in MCA stroke (363) and improve recovery in cerebral damage induced by experimental hematoma (363). However, in gerbil and dog models of stroke, TRH fails to display efficacy (352).

c. TRH and respiration. icv TRH, in doses as low as 3 ng, significantly elevates blood pressure and heart rate in anesthetized rats, but a minimum 16 ng is required to raise the respiratory rate (364). Respiratory frequency is increased much more than tidal volume (16). Increases of respiratory frequency greater than tidal volume are also seen in conscious rats (365) and in rabbits (366). icv TRH raises respiratory rates in anesthetized rats partly through a DA D2 receptor mechanism (332). Further, icv TRH-induced respiratory stimulation is potentiated by pretreatment with naloxone, methylatropine, or low doses of GABA, but is unaffected by β -adrenergic blockade and is independent of TSH (367). TRH antagonism of opiate-induced respiratory depression (368) is described in more detail below in *Subsection 7*.

In isolated brain stem-spinal cord preparations from rat neonates, bath application of TRH induces respiratory rhythmic neural discharges (369). icv TRH produces rhythmic bursting activity in neurons of the NTS (370), and local injection of TRH into the NTS induces tachypnea, although with a slower onset than seen with icv TRH (367). The respiratory effect occurs in the absence of any change in blood pressure or heart rate. Shortening of inspiratory times, but not tachypnea, results from TRH injections into the ROb (371). Tachypnea, without cardiovascular or locomotor effects, is seen with microinjection into the interpeduncular nucleus of the reticular activating system (372).

There is significant anatomical support for a role of TRH in respiratory control. Botzinger neurons in the medulla, which inhibit respiratory motoneurons, and the more caudal ventral respiratory group, form close associations with TRH-immunoreactive boutons (373). These connections appear to be functional, since TRH injected into the pre-Botzinger complex in neonatal rat medullary slices increases respiratory discharge frequency (369). TRH-immunoreactive boutons are also prominent near nucleus ambiguus motoneurons that display rhythmic fluctuations with phrenic nerve discharges (374).

Developmental studies support a direct effect for TRH on hypoglossal motoneurons in the caudal medulla. These neurons innervate tongue muscles critical for airway inspiratory control and display respiratory-related activity. TRH increases hypoglossal neuron discharge frequency, duration, and amplitude in neonatal mouse slices. In adult rat brainstem slice preparations, high doses of TRH depolarize hypoglossal neurons and reduce their firing threshold (375). More rostral to the hypoglossal nucleus, TRH enhances the responsiveness of inspiratory neurons in the ventrolateral medulla (376).

iv Or ic TRH significantly stimulates diaphragmatic activity and antagonizes morphine depression of diaphragmatic activity (377). TRH potentiates the excitability of diaphragmatic motor nerve terminals (378). Further, injections

of TRH as low as 1 ng into the retrotrapezoid nucleus of anesthetized rats increase phrenic nerve firing frequency and amplitude. Only at 5 ng does TRH raise blood pressure. Both CHP and TRH-OH increase phrenic nerve firing frequency, but not amplitude, starting at 5 ng (379).

Preclinical literature describes the importance of thyroid hormones and steroid treatment on fetal lung development (380). However, in human trials TRH coadministration in steroid and surfactant therapy does not reduce newborn respiratory distress syndrome, chronic lung disease or associated neonatal complications, or death (381, 382).

3. *TRH and seizure modulation.* TRH was first reported to potentiate the anticonvulsant actions of phenobarbital in mice in 1975 (383) and has since been shown to be anticonvulsant in multiple animal models of seizures (384–386). Despite extensive preclinical data indicating that TRH is likely to serve as an anticonvulsant, or a potentiator of known anticonvulsants, few large trials with this agent have been conducted. For intractable epilepsy, modest results have been achieved (387, 388).

a. TRH and electroconvulsive seizures: One seizure paradigm commonly used to test the efficacy of anticonvulsant drugs is electroconvulsive seizure (ECS) (389). A single stage 5 seizure, induced after five ECS treatments given on alternate days (ECS \times 5), elevates TRH in hippocampus, amygdala/pyriform cortex, whole cortex, and striatum, 48 h post seizure (17). Subconvulsive shocks given on alternate days for 5 days result in regulation only in the striatum. A separate study reported that ECS \times 5 decreased NAc and lumbar spinal cord TRH 24 h after the last shock (390). In sum, ECS effects center on the hippocampus, amygdala and surrounding cortex, and the dorsal and ventral striatum.

The hippocampus most consistently demonstrates TRH induction after chronic ECS (391). A significant percentage of hippocampal TRH derives from extrinsic sources. ECS induction of prepro-TRH mRNA in the entorhinal cortex presumably leads, via the perforant pathway, to TRH increases in the dentate gyrus (17). In the hilar subregion of the hippocampus, which contains few or no perforant path terminals, fimbriae-fornix lesions do not block ECS induction of TRH, consistent with endogenous biosynthesis (392). There is no difference in basal TRH release from hippocampal slices dissected from ECS-treated vs. sham-treated animals. However, potassium-stimulated TRH release increases linearly 12, 24, and 48 h post seizure, and tissue content remains uniformly elevated throughout the postseizure period (393). Thus, in addition to elevations in steady state levels, there is a time-dependent shift of intracellular TRH into a potassium-responsive pool (394), which may enhance TRH release in response to afferent signaling.

Given the documented effect of TRH to increase cholinergic transmission in the hippocampus (395) and cortex (396, 397), it is logical to speculate that TRH could be used to reverse ECS-induced neurochemical and behavioral deficits (398, 399). Indeed, in rats post-ECS performance deficits are reversed by TRH.

b. TRH in kindled seizures: A second paradigm used to model epilepsy is kindling, where repeated electrical or

chemical stimulation of limbic structures progressively lowers seizure threshold until an initial subthreshold stimulation becomes capable of reliably stimulating generalized (stage 5) seizures (400). In fully amygdala-kindled rats (five consecutive stage 5 seizures), prepro-TRH mRNA levels are significantly elevated 24 h following a stage 5 seizure in the dentate gyrus granular layer and the pyriform, entorhinal, and perirhinal cortices (401, 402). More detailed time course studies in fully kindled animals report significant elevations of prepro-TRH mRNA in the dentate gyrus granular layer, several nuclei of the amygdala, and layers II and III of the pyriform and entorhinal cortices. Increased levels are detected 3 h post seizure, peaks occur at 6–12 h post seizure, and levels return to baseline 24–48 h post-seizure (246, 392). The time course is similar in all regions, although slightly delayed in entorhinal cortex. The induction in prepro-TRH mRNA seen after full kindling may be observed unilaterally 24 h after partial kindled (stage 1–4) seizures, but is reliably and bilaterally observed only after fully kindled, generalized (stage 5) seizures (401).

Kindled seizures induce c-fos mRNA and Fos-related peptides, which in turn are postulated to induce prepro-TRH transcription via the AP-1 site in the prepro-TRH gene promoter (155, 403). Fos-like immunoreactivity and prepro-TRH mRNA are extensively colocalized, in some cases in up to 70% of cells, in the pyriform cortex, entorhinal cortex, and dentate gyrus granule cells (404). A second potential transcriptional regulator of prepro-TRH gene expression, corticosterone, also is rapidly induced during kindled seizures (166, 405).

Carbamazepine given contingently with kindling treatments attenuates prepro-TRH mRNA increases in the dentate gyrus, pyriform cortex, and ipsilateral entorhinal cortex. No attenuation is seen when carbamazepine is given non-contingently, *i.e.*, after the kindling treatment (406). These results are intriguing, indicating that the carbamazepine-TRH interaction might be altered by behavioral or other nonpharmacological interventions.

Regulation of TRH peptide levels, measured 48 h post seizure, correlates with the progression of amygdala kindling. Partial kindling induces TRH in pyriform cortex, with greater regulation in stage 3–4 seizures than stage 2 seizures (17, 407). In fully kindled rats, TRH is increased even more in pyriform cortex 48 h after stage 5 seizure. Increases are also seen in cingulate and frontal cortex, hippocampus, amygdala, and ventral striatum (408, 409). Similarly, chemically kindled rats show TRH induction in hippocampus, amygdala, pyriform cortex, and anterior cortex 48 h after stage 5 seizures (17). Both kindling and TRH regulation after kindling persist for 6 months after kindling.

In fully kindled animals, all subregions of the hippocampus show reduced levels of TRH 1 h after seizure, consistent with synaptic release and rapid degradation. Levels rise to control levels at 6 h, are elevated at 24 and 48 h, and again return to control levels at 144 h after seizure. Increases are largest in the dorsal hippocampus, including the dentate gyrus, hilus, and CA3 region. It is hypothesized (17, 409) that TRH elevations may mediate the postictal refractory period (410). In fully kindled rats, bilateral hippocampal TRH in-

fusions decrease seizure after-discharge duration and overt seizure duration in a dose-dependent manner, consistent with an anticonvulsant action in the hippocampus (394).

TRH receptor binding in the dentate gyrus and perirhinal cortex is decreased in amygdala-kindled rats compared with sham-kindled animals (402, 406). After a single stage 5 seizure in electrically kindled rats, hippocampal membrane TRH receptor binding is reduced 23–29%, and amygdaloid membrane binding is reduced by 21–22% (409, 411). Curiously, in amygdala-kindled rats, dorsal striatal receptor binding is increased 24 h after seizure and persists significantly elevated at 21 days, although no regulation in striatal TRH is reported. Thus, only in certain brain regions do receptor adaptations appear to compensate for elevated TRH levels.

c. TRH in chemically induced seizures: TRH regulation in other types of seizures differs somewhat from that seen in ECS and kindling. Limbic seizures induced by systemic kainic acid substantially increase TRH in the posterior cortex and in the underlying dorsal and ventral hippocampus (412). Smaller increases are detected in the anterior cortex, amygdala/pyriform cortex, and corpus striatum. The increases in TRH are longer lasting than described for ECS, peaking at 2–4 days and resolving by 14 days, except for the dorsal hippocampus, where TRH elevations persist beyond 2 weeks. TRH is rapidly elevated in the septum, hippocampus, and thalamus/midbrain after a single pentylenetetrazol-induced seizure. Pyriform cortex was not tested (384). Pentylenetetrazol-induced tonic-clonic seizures in dogs increase TRH in pyriform and frontal cortex, hippocampus, and amygdala 48 h after seizure (413). Soman-induced seizures mediated by excessive cholinergic activity result in particularly high induction of TRH in frontal cortex, hippocampus, pyriform cortex, and entorhinal cortex, and lower induction in the amygdala (414).

d. Mechanisms of TRH anticonvulsant action: Irrespective of the precise mechanism of seizure induction, repeated seizures ultimately induce prepro-TRH mRNA in pyriform cortex, amygdala, and hippocampus. These areas contain well characterized TRH receptor binding (415, 416). Further, the pyriform cortex region is a primary site for initiation of grand mal seizure activity, and the prepyriform region is exquisitely sensitive to direct application of chemical convulsants (417). One mechanism of TRH anticonvulsant effects may be through inhibition of L-glutamate excitation of neurons (418, 419), especially neurons of the perforant pathways synapsing with dentate gyrus granule cells (420). This inhibition would increase after treatments such as kindling that elevate TRH, providing a feedback control for further sensitization in phenomena such as kindling and long-term potentiation. Knoblich and Kubek (407) suggest that TRH may be coreleased with excitatory neurotransmitters at these sites as a means to modulate neuronal response. If TRH is inhibitory to calcium influx secondary to reducing excitatory amino acid neurotransmission, it may also serve a neuroprotective role. Also it is speculated that hippocampal TRH may interact with coexpressed endogenous opioid peptides in seizure-involved pathways to modulate seizure activity (93).

4. TRH and motor control.

a. TRH stimulates ventral horn motoneurons: Bulbosplinal neurons that express prepro-TRH mRNA and iTRH descend from the medullary raphe nuclei, and the parapyramidal and paraolivary regions, to end in close apposition to motor neurons in lamina IX, and sparsely in lamina VIII, of the ventral horn of the spinal cord (421, 422). The raphe projections provide dense innervation of spinal motoneurons and are likely to enhance motor excitability, principally of proximal muscle groups (423, 424). In rat (425), rabbit (426), and human (427) spinal cord, the highest concentrations of TRH are found in the ventral horn. TRH is present in large granules within terminal boutons that synapse with dendrites, supporting a synaptic role (422). Some 60–90% of these bulbospinal neurons coexpress 5-HT with TRH (428, 429). Immunocytochemical and ablation studies also support coexpression of TRH with SP (422), although SP appears to have a more prominent role in autonomic nervous system function than in voluntary motor control (430). Surprisingly, in most species the ventral horn of the spinal cord is not enriched in TRH receptor binding (431). However, the human spinal cord displays elevated TRH receptor densities in laminae IX, which contains α -motoneurons (432).

An extensive literature describes TRH excitation of spinal cord ventral horn motoneurons (423, 433, 434) and hypoglossal motor neurons (435) by suppression of a distinct K^+ current and development of an associated Ca^{++} -sensitive inward current (436). In addition, TRH enhances motor neuron firing in response to excitatory amino acids (423), increases motoneuron recruitment by antidromic stimulation (437), and depolarizes ventral roots (438). The net effect of this excitation is augmentation of muscle tone, contractility, and spinal reflexivity (438). It should be noted that while TRH, SP, and 5-HT each can enhance excitatory amino acid activation of motoneurons, TRH excitation of ventral horn motoneurons is slower and less reliable than that observed with application of 5-HT or SP (423, 439). Neither TRH-OH nor CHP have any demonstrated effect on spinal motoneurons, so motoneuron effects are presumed to be direct actions of TRH (434, 436).

Denervation of the plantar foot muscles by botulinum toxin injection reveals reinnervation deficiencies in adult rats that have undergone ablation of the descending bulbospinal 5-HT/TRH neurons (440). However, gross motor performance, muscle cell count, electrophysiological properties, or α -motoneuron counts are not made abnormal by this ablation, arguing for an insignificant role for TRH in adult animals. White *et al.* (439) argue that TRH function is more significant in developing animals or on damaged motoneurons. TRH trophic effects (441) and enhancement of contractility (442) are demonstrated best in embryonic/neonatal preparations. More importantly, TRH-induced depolarization shows significantly less tachyphylaxis in isolated neonatal rat spinal cord preparations than in adult preparations (443). NE inputs in the ventral horn enhance both microiontophoretically applied TRH-induced excitation of motoneurons (444) and behavioral excitation elicited by intrathecal

TRH (426). Thus, TRH may function only under certain physiological states, such as stress or healing, that were not well tested in previous paradigms.

In preclinical studies, TRH displays significant beneficial effects in the Rolling mouse Nagoya (RMN), a mouse model of ataxia (445, 446). TRH is also effective in ameliorating ataxia induced by 3-acetylpyridine degeneration of the inferior olive in rats (446) and by cytosine arabinoside treatment in mice (447). Immobility and fall index scores for other ataxic mice models, including *staggerer*, *reeler*, *weaver*, and mice with Purkinje cell degeneration, are reduced by TRH (448, 449). The NMDA antagonist MK-801 completely blocks TRH improvement of ataxia induced by 3-acetylpyridine, providing one of the few clues as to how TRH might mediate its antiataxic effects (450).

Several clinical studies support potential utility for TRH in the treatment of inherited ataxias such as spinocerebellar degeneration. Several studies demonstrate improvements by TRH of motor, oculomotor, and electrophysiological abnormalities in inherited ataxias such as spinocerebellar degeneration (451, 452). For the motor neuron disease amyotrophic lateral sclerosis (ALS), studies using higher doses of TRH demonstrate some transient improvement in symptoms, particularly speech, swallowing, and respiratory function. Unfortunately, longer term results with systemic or intrathecal TRH (453, 454) indicate that TRH fails to slow progressive motor neuron loss and provides only temporary symptomatic relief.

b. TRH promotes recovery in experimental spinal cord and brain injury: TRH accumulates superior to the site of traumatic or ischemic spinal cord injuries (455, 456). In both cervical and lumbar injuries, TRH elevations are accompanied by modest but detectable down-regulation of TRH receptor binding proximal to the injury, in lamina X and the ventral horn gray matter, but not in the dorsal gray (457). The decrease in TRH receptor binding is detectable 48 h after injury and recovers by 3 weeks. Recovery of TRH receptor binding parallels the functional neurological recovery that occurs late after CNS injury (458). In mouse models of spinal cord motoneuron degeneration, TRH metabolism and levels are increased at sites of damage (459, 460).

Whether increased TRH is a neuroprotective adaptation or a damaging mediator of the disease process is unknown. However, it appears likely that TRH is beneficial. TRH is superior to naloxone or high-dose steroid treatment in promoting recovery in experimental CNS trauma models (461). In most studies, sc or iv TRH improves electrophysiological recovery of damaged spinal cord tissue (462, 463). Improvement may result from trophic effects of TRH on spinal motoneurons (204, 464, 465), from the ability of TRH to increase spinal cord blood flow (466, 467), or from the ability to reduce edema (468) at the site of spinal cord injury. TRH efficacy in spinal trauma recovery requires continuous intrathecal infusion of native TRH, presumably due to its short half-life. Further, while TRH and all its analogs activate spinal motoneurons (434), only analogs that preserve the C terminus of native TRH are beneficial in spinal trauma, indicating the healing and activating effects of TRH are separately mediated (469).

TRH and its analogs are also efficacious in promoting recovery in animal models of head injury. Cats with brain stem compression injury show improved neurological and EEG parameters (470), and mice with head impact injuries demonstrate less behavioral disturbance (471), after TRH treatment. In rats with fluid percussion-induced brain injury, the TRH analog NS-3 given 30 min after injury improves survival, neurological parameters, and motor function at 24 h post injury, and improvements persist for at least 4 weeks (472). There is significant interest in TRH as a treatment for human traumatic brain injury and spinal cord injury, either to prevent damage progression or to speed neuronal recovery (473). However, TRH effects on head trauma, or recovery in spinal cord trauma, have been modest at best (474).

5. TRH and antinociception. The presence of TRH and TRH receptors in the midbrain PAG, the raphe nuclei, and, to a limited extent, in the dorsal horn of the spinal cord is highly suggestive of a role in pain modulation (93). Antinociception to chemical stimuli has been demonstrated in mice with iv and sc TRH (475) and for icv TRH against visceral chemical and mechanical pain (476). In the rat, icv TRH increases reaction latencies to visceral acetic acid (477). TRH displays potencies comparable to morphine in some of these studies. Thermal analgesia with icv TRH has been demonstrated in the hot-plate test (476) and less frequently in the tail-flick test (477). icv TRH also potentiates stress-induced analgesia, including foot shock-induced analgesia and swim-induced analgesia (478). In general, the antinociceptive effects of TRH are short lived, typically lasting for less than 15 min (479).

icv CHP in mice is variably reported to induce antinociception to mechanical, thermal, and chemical stimuli (480), or to mechanical stimuli but not chemical stimuli (481). This effect is significantly less potent, but longer lasting, than that of TRH and is antagonized to a greater extent by sc naloxone than is TRH. TRH-OH also is reported to be antinociceptive to mechanical and chemical stimuli in mice (481). Thus, it is likely that both TRH and its metabolites are active in TRH-induced antinociception in its various forms.

The PAG is critical for integration of pain perception and the behavioral response to pain (482). The PAG expresses moderately high levels of prepro-TRH mRNA (28, 249), and significant amounts of iTRH are detected in cell perikarya and neuronal fibers of the PAG (93, 421, 483), and TRH receptor binding is detected at moderate levels (484) throughout the PAG. The neuroanatomical distribution of TRH and TRH receptor binding in all regions of the PAG implicates it in both opiate-dependent and opiate-independent pain mechanisms. The dorsolateral PAG mediates antinociception that is not modified by naloxone and is presumed to be opiate independent, while the ventrolateral PAG is more commonly associated with opioid-dependent antinociception (482). This may relate to the finding that TRH infusions into the ventral PAG decrease cold-water swim-induced antinociception in the rat in a dose-dependent manner, while dorsal PAG infusion has the opposite effect (485). Curiously, TRH in both placements antagonizes morphine antinociception and reduces swim-induced hypothermia and morphine-induced hyperthermia, while the converse, analgesia produced by TRH infusion into the ventral PAG,

is blocked by opiate antagonists (486, 487). TRH has no demonstrable effect on μ -, δ -, or κ -opiate receptor binding, or receptor occupancy, arguing against direct TRH regulation of opioid peptide release. It is postulated that TRH activates one or more types of inhibitory interneuron, which in turn reduce excitation of pain-excited opiate-responsive neurons in the PAG.

Complete blockade of TRH-induced antinociception by opiate antagonists occurs in some PAG infusion paradigms, but not with systemic TRH, which presumably activates non-opiate pain pathways in which TRH functions. Separate studies indicate that TRH injections into the NAc (488) and amygdala (477) produce antinociception. TRH antagonizes NT-induced antinociception (489) and both TRH and CHP antagonize THC-induced antinociception (194).

Clearly then, the relationship of TRH antinociception to endogenous and exogenous opiate systems is complex. As opposed to opiate-induced hypothermia, locomotor depression, catalepsy, and respiratory depression, systemic TRH does not antagonize acute opiate-induced analgesia (490), although hyperalgesia induced by high-dose naloxone is antagonized by TRH (491). In contrast to the case with drug-naive mice, TRH does not display antinociceptive effects to chemical stimuli in morphine-tolerant mice (487). In particular, Bhargava and colleagues (490) argue that TRH interacts preferentially with the κ -opiate system in mice. Other studies (487) indicate that tolerance either to morphine (μ - and δ -specific) or ethylketocyclazocine methanesulfonate (κ -preferring) equally antagonize TRH antinociception. Most studies indicate that TRH-opiate interactions do not occur via TRH modulation of opiate receptor binding, or through TRH-stimulated release of endogenous opioids. Rather, TRH acts through intermediary systems that, in turn, modulate opiate-mediated pain transmission, and perhaps vice versa. In the spinal cord, likely intermediary systems involve 5-HT and SP (492).

PAG-mediated antinociception is mediated via outputs to the ROb, RPa, and RMg (482), nuclei that express high levels of prepro-TRH mRNA (249). Further, antinociception by excitation of the nucleus reticularis paragiganticularis (RPGi) is mediated, in large part, through reciprocal connectivity to the RMg (493). TRH into either the RMg or RPGi is antinociceptive (477). icv TRH also inhibits pain-excited neurons in the mesencephalic reticular formation (MAF) (494). The MAF is believed to form a second integration circuit for autonomic response to painful stimuli (495).

In addition to the descending bulbospinal pathway, TRH is also found in an intrinsic system of cell bodies in laminae II and the lamina II/III border of the dorsal horn (251, 496). Dorsal horn iTRH is not depleted by 5-HT neurotoxins, unlike ventral horn TRH (250). Laminae II also exhibit a high level of TRH binding in many species (250, 432, 484). Intrathecal TRH antagonizes morphine analgesia in the tail-flick test at most doses, indicating that the net action of TRH within the dorsal horn is to enhance transmission of nociceptive somatosensory information (492). iv TRH is reported to facilitate nociceptive transmission through the dorsal horn via positive modulation of NMDA receptor-mediated transmission (497).

6. *TRH in thermoregulation.* TRH plays a prominent role in integrating a number of thermogenic responses to cold (509). CNS injection of TRH elevates body temperature (498), and TRH antagonizes the hypothermic effects of a number of agents, including barbiturates, ethanol, chlorpromazine, bombesin, NT, and β -endorphin (204). Systemic TRH antagonizes morphine-induced hypothermia, while having little effect on analgesia (499). icv Anti-TRH antibodies in rats produce hypothermia, supporting an endogenous role for TRH in body temperature elevation (500). A principal site of TRH thermoregulation is the anterior hypothalamic POA (498). TRH into the POA inhibits heat-sensitive neurons and activates cold-sensitive neurons (501), which results in increased body temperature through peripheral vasoconstriction, increased metabolic heat production, and shivering (502). These effects require intact catecholamine neurotransmission (502).

Ablation of the POA does not eliminate TRH antagonism of pentobarbital-induced hypothermia, indicating that sites other than the POA can mediate TRH thermoregulation (503). Cold exposure elevates prepro-TRH mRNA levels (504) and TRH secretion (505) in the PVN. These changes elevate thyroid hormones and increase heat generation in brown adipose tissue (506). Systemic TRH has similar effects, e.g., systemic TRH improves thermoregulation in neonatal lambs through increased fat oxidation (507). We note that cold exposure elevates TSH levels before TRH levels, probably because SRIF, which tonically inhibits TSH secretion, is rapidly down-regulated in the PVN by cold (508).

Cold-induced increases in prepro-TRH mRNA also are seen in the DMN (509) and caudal raphe nuclei (510). The raphe nuclei, which receive sensory information from the skin, project to spinal cord preganglionic sympathetic neurons. Further, the raphe nuclei provide TRH afferents to the NTS (511), which, in turn, projects to the DMN, and then the spinal cord preganglionic neurons. 5-HT projections from the NTS to the PVN provide feedback regulation to this stimulation (512). Excitation of spinal cord preganglionic sympathetic neurons results in postganglionic NE release and increased facultative thermogenesis via β - and α 1-adrenoreceptors on brown adipocytes (513). Additional stimulation comes from direct projections of TRH neurons from the dorsal cap of the PVN to preganglionic sympathetic neurons in the thoracic and sacral spinal cord (514).

In mice, icv TRH and CHP antagonize ip THC-induced hypothermia (517). However, CHP elicits hypothermia when injected into the cerebral ventricles, an effect antagonized by TRH (515). The POA is believed to be the sole site mediating this action of CHP (516). Thus, the hypothermic response to icv TRH seen under certain conditions such as warm environments may result from TRH catabolism to CHP (501).

7. *TRH and drugs of abuse.* The psychomotor theory of addiction states that a common biological mechanism mediates both positive reinforcement and motor activation by drugs of abuse (518). Given the ergotrophic effect of TRH as a locomotor activator, interaction between TRH and drugs of abuse is likely. While the main focus of drug abuse research has centered on two principal loci in the mesolimbic DA pathway, the NAc and the VTA (519), a more extensive network,

the "limbic-motor circuit," with inputs to the NAc coming from many limbic areas and outputs going to both limbic areas and motor areas, is now appreciated (520).

In rats, systemic TRH and intra-NAc TRH mimic cocaine by inducing locomotor activation via release of DA and 5-HT in the NAc and striatum (222, 521, 522). Conversely, DA D2 agonists increase TRH release from striatal and NAc slices (523). Downstream components of the HPT axis act to reinforce psychostimulant effects (524). TRH neurons of the PVN receive DA and NE inputs that are regulated by cocaine (124). Further, DA and cocaine both activate the HPA axis; stress (or CRF) and cocaine elevate NAc and medial prefrontal cortex DA and cause similar neuronal adaptations (525, 526).

Few studies have directly examined the link between TRH and psychostimulants. Acute amphetamine lowers TRH in the caudate, NAc, and lateral septum (527, 528). Over time, TRH levels show some adaptation to chronic amphetamine, and TRH receptor binding increases (528). Acute cocaine significantly decreases prepro-TRH mRNA levels in the amygdala and hippocampus, 45 min after injection (529). Chronic cocaine regulates prepro-TRH mRNA in the NAc, amygdala, hippocampus, and hypothalamus. Prepro-TRH mRNA regulation is strongly dependent on the length of time after cocaine cessation and persists beyond 72 h post injection in the amygdala.

The role of TRH in morphine actions also is not well understood. TRH antagonizes a number of morphine's depressant effects, including sedation, hypothermia, and catalepsy (530, 531). Chronic TRH inhibits the development of tolerance to opiate-induced hypothermia and catalepsy (499, 532). δ - and κ -opiate receptor activation reduces TRH receptor binding, but TRH does not effect opiate binding (533). More directly, morphine reduces cortical and diencephalic TRH concentrations (534). Within the HPT axis, a clearer relationship between TRH and opioids exists. Morphine and opioid peptides reduce plasma TSH (535) and blunt cold-induced TSH release (536). Further, exogenous morphine at pharmacological doses inhibits TRH release via opiate receptors on TRH-secreting hypothalamic nerve terminals (509).

TRH is more strongly implicated in opiate withdrawal. While cessation of chronic cocaine use induces relatively little physical withdrawal (537), chronic morphine results in the development of physical dependence and the aversive state of withdrawal upon cessation of morphine use (537). The expression of the physical symptoms of withdrawal is mediated principally by the LC and PAG (538). A large body of data indicates that the intrinsic NE neurons of the LC undergo an up-regulation of their cAMP second messenger system in response to chronic morphine. When unopposed during morphine withdrawal, the up-regulated cAMP system drives increased firing by LC neurons (539). Extrinsic excitatory amino acid inputs from the nucleus paragigantocellularis lateralis contribute an additional 50% to LC neuronal excitability during withdrawal (540, 541). The medial hypothalamus, medial thalamus, amygdala, frontal cortex, hippocampus, and RMg also are implicated in withdrawal (542, 543).

The PAG expresses high levels of prepro-TRH mRNA (28). Mature TRH (483) and TRH receptor binding (484) are

present in moderate levels throughout the PAG. Prepro-TRH mRNA is strongly induced in the PAG during opiate withdrawal (28). Fos-like immunoreactivity is greatly increased in the ventrolateral PAG during withdrawal (544) and may mediate induction of prepro-TRH mRNA (401).

While TRH levels in the PAG remain unchanged during opiate withdrawal (154), it has been found that ic TRH prevents withdrawal-induced hypothermia and decreases jumping during withdrawal in morphine-dependent mice (545). In contrast, ic TRH induces wet-dog shakes in normal animals, arguing that it augments withdrawal-like symptomatology (546). Opiate withdrawal increases TRH in the lateral hypothalamus, suggesting this region may also play a physiological role in opiate withdrawal (154). Thus, much remains to be learned about TRH effects on opiate withdrawal at sites other than the PAG.

The TRH analog TA0910 reduces alcohol-intake in alcohol-preferring rats (547) and in primates (548) in a dose-dependent manner. This appears mediated by DA D2 receptors (549). Behavioral reward to alcohol, as measured by punished responding rates, is enhanced by iv TRH (550). Alcohol alters TRH receptor binding (551). In long-sleep (LS) and short-sleep (SS) mice that display differential CNS sensitivity to ethanol, SS mice have greater sensitivity to TRH than LS mice during postnatal days 8–14 (552). It is hypothesized that a TRH receptor-mediated alteration results in enhanced development of the thyroid gland in SS mice. Alcohol-preferring rats compared with nonpreferring rats have significantly lower TRH levels in the medial and lateral septum. Upon exposure to alcohol, preferring rats are able to right themselves earlier than nonpreferring rats, and this correlates with elevations of medial septal TRH. However, these findings may be nonspecifically related to the analeptic action of TRH in this region (202). Of unclear relationship to alcohol preference, chronic ethanol in rats partially "uncouples" PVN TRH expression from peripheral thyroid response (553, 554) and, like opiates, ethanol blocks the TSH response to cold (553, 554).

A wide range of other addictive substances alter TRH receptor binding, including THC and chlordiazepoxide (551, 555). Furthermore, as described above, behavioral reward to pentobarbital and chlordiazepoxide, as well as alcohol, as measured by punished responding rates, is enhanced up to 3.5-fold by iv TRH (550). Goeders *et al.* (556) recently presented evidence that levels of benzodiazepine receptor binding are affected by the development of either behavioral tolerance or sensitization to cocaine. Conversely, benzodiazepines specifically attenuate cocaine self-administration (557). Since a number of benzodiazepines displace ^3H -methyl-TRH from TRH receptors (558, 559), this provides another suggestive link between TRH and drugs of abuse.

In summary, there are multiple pathways through which TRH affects virtually all classes of abused drugs. Understanding these interactions is likely to advance our understanding of addiction in general. More importantly, this understanding may provide new pharmacological approaches for the clinical treatment of substance abuse.

8. *TRH outside of the CNS.* TRH is phylogenetically old, present in invertebrates such as the lamprey that lack TSH or

the snail that lacks a pituitary (560, 561). It appears that mammalian endocrine functions for TRH have been "co-opted" (562) for a peptide already functioning in more basic ways. TRH is detected in many nonneural vertebrate tissues, although its functions in these tissues are not well understood. In most cases, peripheral TRH is not regulated coordinately with the HPT axis (563).

Prominent among the TRH-containing tissues are the gastrointestinal organs, including the stomach, duodenum, small intestine, colon, and rectum (562), where TRH may have peripheral effects to modulate gastrointestinal contractility. The pancreas is a rich source of TRH. Indeed, in neonates a significant portion of circulating TRH is derived from pancreas (561). Prepro-TRH mRNA is expressed in β -cells of the pancreatic islet (564), and TRH and somatostatin have opposing paracrine effects on glucagon secretion (565).

TRH is present at high levels in the genitourinary system including the ventral prostate, Leydig cells of the testes, the epididymis, and seminal vesicles (566, 567). Interestingly, propylthiouracil-induced hypothyroidism increases TRH in prostate and testis but reduces TRH in epididymis (566, 568). TRH receptor mRNA is expressed in the ovary and uterus (569), and TRH is present in placenta, amniotic fluid, and breast milk (177, 570). Again, reproductive TRH may act as a paracrine regulator (571).

TRH is also present in retina (572, 573), where its levels are light entrained (572). icv TRH raises intraocular pressure and induces marked mydriasis via combined sympathetic and parasympathetic effects (574). TRH receptor mRNA is found in human peripheral blood monocytes (PBMCs) and rat splenocytes (575). It affects secretion of TSH and immunoglobulins from blood elements (576, 577) and may be a trophic factor for certain blood elements (578). In the heart, TRH is expressed and has direct ionotropic effects (579, 580).

IV. Function of non-TRH pro-TRH-Derived Peptides

The rat TRH precursor yields, in addition to TRH, seven pro-TRH-derived peptides that display region-specific distributions (19, 82, 87, 96, 581). In addition, TRH progenitors, *i.e.*, extended forms of TRH that have not been fully processed, are found. Of these, the following discussion will focus on the immediate TRH progenitor, TRH-Gly. From the first immunohistochemical studies describing the distribution of pro-TRH, it was clear that precursor immunoreactivity was present in areas of the brain where mature TRH had not been localized (82, 95, 582). These regions included nuclei in the olfactory bulb, sexual dimorphic area of the POA, reticular nucleus of the thalamus, amygdala, hippocampus, cerebral cortex, and PAG (95). These conclusions were based on immunohistochemical localization of TRH in colchicine-treated animals. However, TRH localization based on micropunch isolation and RIA, where animals need not be colchicine treated, has detected TRH in many of these regions, including the amygdala and surrounding cortical regions, the hippocampus, and PAG. Still, certain regions, such as the reticular nucleus of the thalamus, appear to contain little or no mature TRH, while studies using antisera

raised against non-TRH pro-TRH-derived in brain extracts demonstrate significant amounts of pro-TRH-derived products in this nucleus (Table 1). Thus, while *in situ* immunohistochemical studies remain of value, one must be aware that they are beset with difficulties in variable immunoreactivity in different loci of the brain, and they lack precise definition of the immunoreactive peptide moieties that are being detected.

In comparison to the known roles of TRH reviewed in Section III.B, there is precious little known about the biological activities of the other pro-TRH-derived peptides. In the hypothalamus and testis, both these non-TRH peptides and TRH are regulated within HPT parameters, while in other tissues, both TRH and non-TRH peptides are not (563, 583). Both TRH-Gly and prepro-TRH₁₇₈₋₁₉₉ are regulated by dexamethasone in the hypothalamus, but not in cerebellum, brain stem, retina, and stomach (583). Thus, clues to the roles of pro-TRH-derived peptides other than TRH must come from an examination of their regional distribution or evidence of regulation under specific physiological or pathological conditions.

A. prepro-TRH₁₆₀₋₁₆₉ (pST₁₀)

Prepro-TRH₁₆₀₋₁₆₉ is the best characterized of the non-TRH pro-TRH-derived peptides. It is released from rat hypothalamic slices and the ME, thus making a hypophysiotropic role likely (20). Prepro-TRH₁₆₀₋₁₆₉ (also known as Ps4 and TRH-potentiating peptide) enhances TRH-stimulated TSH release from the anterior pituitary and stimulates TSH β gene promoter activity (21). Thus, it acts in an opposite manner to feedback by T₃, which decreases TSH secretion (172) and inhibits TSH subunit gene expression (584). The peptide has been isolated from bovine hypothalamus and its amino acid sequence confirmed by Edman degradation (585). Prepro-TRH₁₆₀₋₁₆₉ is also unique in that a receptor for this peptide has been characterized (see Section VI) (586). Prepro-TRH₁₆₀₋₁₆₉ receptor binding is developmentally regulated, with an increase from birth to weaning, and then a gradual decline to adult levels at postnatal day 60 (587).

Within the CNS, prepro-TRH₁₆₀₋₁₆₉ is most enriched in hypothalamus, with lesser amounts in the spinal cord and olfactory bulb. The pituitary and striatum contain moderate levels. Its receptor binding is highest in the pituitary, hypothalamus, spinal cord, olfactory bulb, and hippocampus (588). Prepro-TRH₁₆₀₋₁₆₉ is rich in rat testis, but trace levels are detected in urinary bladder, vas deferens, and heart. Receptor binding is high in urinary bladder and vas deferens, heart, and testis (588). In adrenal extracts, RIA detects lesser amounts of prepro-TRH₁₆₀₋₁₆₉, which has been confirmed further in this tissue by chromatographic fractionation. The pancreas contains prepro-TRH₁₆₀₋₁₆₉ within β -cell secretory granules (318). Finally, prepro-TRH₁₆₀₋₁₆₉ function is not restricted to a hypophysiotropic one. In parallel fashion to its role in the pituitary, prepro-TRH₁₆₀₋₁₆₉ does not influence basal gastric acid secretion when injected into the DMN, but does potentiate the ability of TRH to do so. Prepro-TRH₁₇₈₋₁₉₉ has no effect when coinjected with TRH into the DMN, and neither does prepro-TRH₁₆₀₋₁₆₉ when coinjected with TRH into the nucleus ambiguus (22).

ECS elevates prepro-TRH₁₆₀₋₁₆₉ in hippocampus, amygdala, pyriform cortex, and anterior cortex, but not corpus striatum, motor cortex, LC, or ventrolateral medulla (589). In these studies, elevations of prepro-TRH₁₆₀₋₁₆₉ correlated with elevations in TRH-Gly and TRH in the hippocampus, amygdala, and pyriform cortex. Thus, prepro-TRH₁₆₀₋₁₆₉ may share a role with TRH in seizure modulation (17). More recently it has been reported that prepro-TRH₁₆₀₋₁₆₉ levels in the hippocampus and amygdala correlate with immobility times in the Porsolt forced-swim test, leading to speculation that the peptide may act independently or in concert with TRH to affect mood, learning, or memory (589). A review of the distribution and postulated functions of this peptide recently has been presented (590).

B. prepro-TRH₁₇₈₋₁₉₉ (pFE₂₂)

The second most studied non-TRH pro-TRH-derived peptide, prepro-TRH₁₇₈₋₁₉₉, is also released from rat hypothalamic slices and the ME (20, 591) and is localized in dense-core granules of PVN neurons that project down to the ME. prepro-TRH₁₇₈₋₁₉₉ has been reported to be a corticotropin-inhibiting factor, acting to reduce POMC mRNA and inhibit ACTH release (592, 593). These data fit the clinical phenomenon of increased TSH in isolated ACTH deficiency (594). Further, α 2-adrenergic inputs stimulate release of both pro-TRH products and ACTH (595), and thus, prepro-TRH₁₇₈₋₁₉₉ and ACTH may be coreleased into the ME, allowing the pro-TRH product to modulate ACTH release. Despite this logic, the relationship between the HPA and HPT axes remains incompletely understood, and other investigators have been unable to reproduce corticotropin inhibition by prepro-TRH₁₇₈₋₁₉₉ (25). The pancreas produces prepro-TRH₁₇₈₋₁₉₉, within the β -cell secretory granules (316), suggesting a potential involvement for this peptide in the regulation of glucose metabolism. Finally, prepro-TRH₁₇₈₋₁₉₉ acts as a PRL secretagogue in primary pituitary cultures. However, as described in the next section, it may be broken down to peptide products that also induce PRL secretion, so a direct effect remains to be proven. Prepro-TRH₁₇₈₋₁₉₉ levels rise during the early phases of suckling in rat pups (91). Again, a precise role for the peptide in lactation, suckling, etc. remains to be determined.

C. prepro-TRH₁₇₈₋₁₈₅ and prepro-TRH₁₈₆₋₁₉₉ (pFQ₇ and pSE₁₄)

Suckling increases prepro-TRH mRNA in PVN and markedly increases TRH release during the first period of lactation (596). In experiments where we coexpressed rat prepro-TRH cDNA with PC1, PC2, and 7B2 in GH₄C₁ cells (Section II.D), we detected two novel peptides, prepro-TRH₁₇₈₋₁₈₅ (pFQ₇) and prepro-TRH₁₈₆₋₁₉₉ (pSE₁₄). These peptides are generated by cleavage of prepro-TRH₁₇₈₋₁₉₉ (pFE₂₂) by PC2 (see Section II.D and Figs. 1 and 6). We subsequently determined that these peptides are present in the rat PVN. In examining whether pro-TRH processing is altered by suckling, we found that in addition to prepro-TRH₁₇₈₋₁₉₉, the products prepro-TRH₁₇₈₋₁₈₅ and prepro-TRH₁₈₆₋₁₉₉ also increase release of PRL from primary pituitary cultures. Prepro-

TRH₁₇₈₋₁₈₅ was the most active PRL secretagogue. In suckling experiments, where pups were separated from their mothers for 6 h and then reunited for 45 min to suckle, a 5-fold increase in PVN prepro-TRH₁₇₈₋₁₉₉ and prepro-TRH₁₈₆₋₁₉₉ and a 6-fold increase in serum PRL were observed over nonsuckling controls. While these data implicate these novel peptides in suckling, or a response to PRL, further experiments are required to rule out nonspecific effects of stress (91).

D. prepro-TRH₅₃₋₇₄ (pFT₂₂)

Prepro-TRH₅₃₋₇₄ displays a unique localization in the rostral two-thirds of the ventrolateral PAG (95, 597). Electrical stimulation of, or injection of excitatory amino acids into, this region of the PAG produces analgesia (598, 599), and this region of the PAG is most sensitive to production of morphine-induced antinociception (600). Although the lateral PAG is more commonly associated with non-opioid-mediated antinociception (482), there is limited evidence that non-opioid antinociception is also mediated within the ventrolateral PAG (601). Thus, prepro-TRH₅₃₋₇₄ may be a candidate molecule to mediate nonopiate pain perception in this region or play a modulatory role in opiate-mediated pain mechanisms.

As described in Section III.B.5, the expression of prepro-TRH mRNA in the reticular nucleus of the thalamus, the PAG, the raphe nuclei, and, to a limited extent, in the dorsal horn of the spinal cord is highly suggestive of a role in pain modulation (93). Prepro-TRH₅₃₋₇₄ similarly is prominent in the ventrolateral PAG, RMg, and thalamic reticular nucleus, suggesting a role in opiate-dependent pain perception (95, 597). The reticular nucleus serves in gating of peripheral somatic sensory information from the dorsal thalamus to the cortex (602). Opiate withdrawal induces prepro-TRH₅₃₋₇₄ in the rat PAG, while TRH levels are unaltered (27), suggesting that the peptide may interact with opiate pain mechanisms in these pathways (406, 597).

E. prepro-TRH₈₃₋₁₀₆ (pEH₂₄) and prepro-TRH₂₀₈₋₂₅₅

As described above (Section III.B.10), during opiate withdrawal prepro-TRH mRNA is increased in the PAG (27, 28). Peptide analysis in the PAG demonstrates an accumulation of the N-terminal peptides prepro-TRH₅₃₋₇₄ (see subsection 7 above) and prepro-TRH₈₃₋₁₀₆, a reduction in the C-terminal peptide prepro-TRH₂₀₈₋₂₅₅, and no change in prepro-TRH₁₇₈₋₁₉₉ or TRH, in opiate-withdrawal *vs.* control animals (154). Opiate withdrawal also increases prepro-TRH₈₃₋₁₀₆ in the lateral hypothalamus. We speculate that during opiate withdrawal, pro-TRH processing may be altered in several brain regions, resulting in increased levels of N-terminally derived peptides (prepro-TRH₅₃₋₇₄ and prepro-TRH₈₃₋₁₀₆) and decreased levels of some C-terminally derived peptides (prepro-TRH₂₀₈₋₂₅₅).

F. TRH-Gly

Immunoreactivity of the immediate precursor of TRH, TRH-Gly, is widespread and detectable throughout the CNS in a similar distribution to TRH and is also present in the

prostate, serum, spleen, adrenals, kidney, and gastrointestinal organs. Levels of TRH-Gly are up-regulated by hypothyroidism and by thermal stress, following the pattern for TRH itself (603). The pancreas contains TRH-Gly within β -cell secretory granules (603). The ratio of TRH/TRH-Gly is highest in pituitary and hypothalamus and much lower elsewhere in neural tissue. TRH-Gly is much better characterized than other pro-TRH-derived peptides discussed above. TRH-Gly is increased in several limbic regions after chronic ECS, including hippocampus and pyriform cortex (391). These authors hypothesize that the increase correlates with effectiveness in increasing swim times in the forced-swim test, and thus it may serve an antidepressant and/or anticonvulsant function. TRH-Gly, independent of conversion to TRH, stimulates gastric acid secretion in a dose-dependent manner, although with a potency 100-fold less than TRH (18). Interestingly, TRH-Gly can directly activate TRH receptors in high concentrations (100 and 1000 nM). The IC_{50} of TRH-Gly for displacement of MeTRH (12 μ M) is significantly higher than the TRH receptor dissociation constant (K_d) of 1.7 nM for MeTRH (604).

TRH in pathological conditions may induce secretion of GH (124). In normal controls TRH-Gly does not increase secretion of TSH, PRL, or GH (605). However, iv TRH-Gly can induce secretion of GH in patients with acromegaly and stimulate PRL and TSH release in women with anorexia nervosa. Preclinical studies indicate estrogen/progesterone treatment, as well as starvation, can enhance the ability of TRH-Gly to stimulate TSH and PRL release, although at a potency far below that for TRH (606). In most cases, TRH-Gly effects are difficult to separate from effects resulting from subsequent conversion to TRH. However, TRH-Gly may prove useful as a pharmacological challenge agent, even if an endogenous role in pituitary release is not confirmed.

V. Non-TRH pro-TRH-Derived Peptides Outside of the CNS

In vertebrates, non-TRH pro-TRH-derived peptides are only reported in tissues where TRH is also found. Their functions are a matter of conjecture, based on their regional and developmental patterns of expression. Anti-pCC₁₀, which detects immediate TRH precursors flanked by paired basic residues as well as larger intermediates, detects immunoreactivity in the pancreas (607), specifically the developing islet β -cells in rat neonatal pancreas (608). Levels of TRH peak at postnatal days 2–4 in parallel with levels of prepro-TRH_{160–169} and prepro-TRH_{178–199}. Both TRH and the pro-TRH-derived peptides are nearly completely abolished by streptozocin treatment, indicating that they are derived principally from β -cells (564). Likewise, prepro-TRH_{53–74}, extended forms of this peptide, and the octa-TRH progenitor are localized to developing β -cells (608, 609). Immunoreactive pCC₁₀ is also found in the retina (607). In adrenal extracts, not only TRH and immunoreactive pCC₁₀ (607), but also prepro-TRH_{160–169}, is detected by RIA and confirmed chromatographically.

In CA77 thyroid parafollicular cells, prepro-TRH mRNA, 7-kDa and 3-kDa species recognized by anti-prepro-

TRH_{53–74} and TRH are detected (86). In thyroid tissue, prepro-TRH mRNA, the 7-kDa and 3-kDa species of immunoreactive prepro-TRH_{53–74}, and immunoreactive prepro-TRH_{115–151} are found. Part of this immunoreactivity comigrates with synthetic prepro-TRH_{115–151} standard on gel filtration and reversed-phase HPLC. In addition, immunohistochemical studies localize prepro-TRH_{53–74} to parafollicular cells in thyroid tissue (610). Thus, within the thyroid, significant biological functions for non-TRH peptides remain to be deciphered.

TRH-Gly is present in high levels in the ventral prostate of the rat, as well as in the testis, epididymis, and seminal vesicles (566). Both TRH and TRH-Gly in the epididymis and prostate are regulated along with the HPT axis under certain conditions (566). Prepro-TRH_{160–169} is also detected in high levels in the testis. Further, prepro-TRH_{160–169} receptor binding is high in urogenital organs, second only to CNS tissues (586). Human placenta contains appreciable quantities of the human TRH progenitor octa-TRH and a larger non-TRH peptide, human prepro-TRH_{192–222} (611).

VI. TRH and Other pro-TRH-Derived Peptide Receptors

A. The TRH receptor

Although a complete discussion of TRH receptors is beyond the scope of this review, some review is important as regards the fate of pro-TRH products that are released. Full-length cDNA clones for the first TRH receptor (TRH-R1) have been identified in mouse (612), rat (613), and human (614). Although there are splice variants for rat and mouse that differ at their C terminus, functional differences between these isoforms, or characterization of their relative occurrence in tissues and under various physiological conditions, has not been determined (615). A second TRH receptor, TRH-R2, has recently been cloned (616, 617). This G protein-coupled receptor is approximately 50% related to the TRH-R1 at the amino acid level and modulates calcium influx upon binding TRH. The expression of TRH-R2 differs from that of TRH-R1, being restricted to the CNS and being enriched in the spinothalamic tract, dorsal horn of the spinal cord, pontine nuclei, and cerebellum. This distribution in the first two sites is suggestive of a role in sensory perception and antinoception, while the later two areas of expression suggest a role in motor control. The identification of TRH receptor species, be they subtypes or members of a new family with distinct pharmacological and/or neuroanatomical profiles, is critical if we are to utilize our growing knowledge of TRH functions in the development of clinically useful therapeutics.

TRH-R1 is highly conserved between species, *e.g.*, at the nucleotide level the human receptor is 90.3% and 89.2% homologous to the mouse and rat receptors, respectively; the three receptors are approximately 95% conserved at the amino acid level. The receptors are members of the G protein-coupled receptor superfamily. Intracellular signal transduction is principally mediated by coupling to G_q and G₁₁. Ligand binding results in activation of phosphoinositide-specific phospholipase C (PPI-PLC) (618), resulting in PIP₂

hydrolysis, and subsequent production of inositol 1,4,5-triphosphosphate (IP3) and 1,2-diacylglycerol. This stimulates increased intracellular calcium, although the exact contributions of increased calcium influx *vs.* mobilization of intracellular stores is controversial (619). Downstream activation of protein kinase C (620), calcium/calmodulin-dependent protein kinase (621, 622), and mitogen-activated protein kinase (623, 624) also occurs. Under specific conditions in certain cell types, the TRH receptor also couples G_{12} and G_{13} and to a G_s -like protein that does not activate adenylate cyclase (625). These complexes are less well studied and represent the minority of TRH signal transduction.

Of particular interest to the potential clinical usefulness of TRH or TRH analogs is the phenomenon of TRH receptor desensitization. Pituitary TRH receptors after several hours exposure to TRH display markedly reduced TSH, but not PRL release. The IP3 response to TRH displays homologous desensitization in as little as 10 sec of TRH exposure in transfected HEK 293 cells (626). This occurs by rapid uncoupling of the receptor and a decrease in PPI-PLC activity. In the same model system, intracellular calcium mobilization displays heterologous acute desensitization, with effects on other receptors whose signal transduction also depends on calcium elevation (627). While most G protein-coupled receptors undergo acute desensitization by phosphorylation (628), this has not been demonstrated for TRH receptors. Neither a specific protein kinase nor calcium concentration have been clearly implicated in TRH receptor desensitization (625). Because G_{11} , G_q , and PPI-PLC have not been shown to be targets for desensitization, it still is believed some form of modification of the TRH receptor may explain acute desensitization. Not to be overlooked, if TRH exhibits slowed dissociation from the TRH-receptor complex, reactivation of the receptor cannot occur (629). Acute desensitization of the TRH receptor is also dependent on cell type, with pituitary cells displaying the most desensitization (630).

A second principal mechanism for TRH receptor desensitization is agonist-induced internalization (631). Up to 80% of TRH receptors are internalized by pituitary cells, with a half-time of 2–3 min (631). Thus, this mechanism is used by TRH receptors to a greater extent than for many other G protein-coupled receptors and may represent a significant mechanism for clearance of secreted TRH. However, recent studies indicate that desensitization does not depend upon internalization, *i.e.*, if internalization is blocked, receptor uncoupling can still mediate desensitization (629). This is similar to the angiotensin II and muscarinic M3 receptors that are also coupled to G_q and G_{11} . While the TRH receptor undergoes internalization and recyclicalization without ligand binding, this “housekeeping” function is slow relative to TRH-induced endocytosis (632). TRH receptor-ligand complexes are internalized in clathrin-coated vesicles (632). A portion of the receptor is targeted to lysosomes, while the remainder is recycled to the cell surface. Similarly, the ligand may remain associated with the receptor to return to the cell surface, or if it dissociates intracellularly, will be degraded in lysosomes or, possibly, reach the cell surface as well (633). Internalization of the TRH receptor is dependent upon sequence motifs within its C terminus (634), as well as sequences within the second transmembrane region or third

intracellular loop that are necessary for G protein coupling (633). Further, optimal rates of internalization appear to require coupling to $G_{q/11}$ and PPI-PLC (635).

Over longer periods of time, TRH receptor binding is also reduced by down-regulation of receptor number. TRH receptor down-regulation occurs in response to TRH, thyroid hormones, and agents that raise cAMP levels (625, 636). Like acute desensitization, receptor down-regulation is dependent upon the cell of expression, as determined in transfected cell experiments (637). The best characterized mechanism of reduced TRH receptor number is reduction in TRH-R1 mRNA levels in GH3 and GH4C1 cells treated with TRH (638, 639). TRH-R1 mRNA regulation is tightly coupled to activation of the TRH signal transduction elements, protein kinase C, IP3, and intracellular calcium (638–640). Desensitization of TRH-R1 by cAMP is also mediated by protein kinase A. Conversely, a number of conditions, including hypothyroidism (641), and treatment with dexamethasone (639), estradiol (642), and cycloheximide elevate TRH-R1 mRNA. Both for TRH-R1 mRNA down- and up-regulation, direct effects on gene transcription rates can be demonstrated.

There is also considerable evidence for regulation of TRH receptor mRNA stability. In GH3 cells transfected with the TRH-R1 coding sequence under control of a cytomegalovirus promoter, TRH increases degradation of TRH-R1 mRNA (643). TRH-R1 mRNA degradation is induced in pituitary cell types by phorbol esters (644). Narayanan and co-workers (645) present evidence in GH₃ pituitary cells that TRH-R1 mRNA degradation is controlled both by *cis*-acting elements within the mRNA 3'-untranslated region and by induction of RNases. Cell transfection studies indicate that regulation of TRH-R1 mRNA degradation does not occur in nonpituitary cells (644, 646). Estradiol-induced increases in TRH receptor mRNA also appear to be mediated through reduction in TRH receptor mRNA degradation (642).

B. The prepro-TRH_{160–169} (pST₁₀) receptor

Of the other pro-TRH-derived peptides, only prepro-TRH_{160–169} has characterized receptor binding (647). Receptors for prepro-TRH_{160–169} seem to be of a single class, with a higher affinity for [³H] pST₁₀ (IC₅₀ = 8.3 ± 1.2 nM) than the native pST₁₀ (IC₅₀ = 9.3 ± 1.2 μM). Recent studies indicate that pST₁₀ receptors cosegregate with S-100 protein-positive cells in pituitary cultures, supporting their expression in the folliculo-stellate cells of the anterior pituitary. Binding sites for pST₁₀ are developmentally regulated, with an increase from birth to weaning, and then a gradual decline to adult levels at postnatal day 60 (587). Signal transduction by these receptors is not yet characterized. Within the CNS, pST₁₀ receptor binding is highest in pituitary, the hypothalamus, spinal cord, and olfactory bulb, as well as the hippocampus (648). Its receptor binding is very high, two-thirds of that in the pituitary, in urinary bladder, and vas deferens and in the heart and testis, at a level equivalent to the hypothalamus (588).

VII. TRH Degradation

The rapid degradation of TRH after release from cells represents a significant drawback in its potential use as a therapeutic agent. To overcome these difficulties, various analogs have been synthesized and evaluated by many groups of investigators (181). Specific enzymes that act on TRH are found in many tissues including the brain, spinal cord, pituitary, liver, kidney, pancreas, adrenal glands, and blood. These enzymes not only inactivate TRH, but act in concert with biosynthetic processes to determine the steady state levels of TRH, and its metabolites, CHP and TRH-OH. The distribution and biological effects of TRH metabolites are discussed more fully in *Section III.B*.

Four key enzymes breakdown TRH: PAP I, PAP II, and thyroliberinase give rise to the stable cyclized metabolite CHP (also known as histidyl-proline-diketopiperazine or His-Pro-DKP), and prolyl endopeptidase gives rise to the deamidated free acid, TRH-OH (182). These enzyme pathways are shown schematically in Fig. 10. In the CNS, the soluble PAP I and prolyl endopeptidase, and the membrane-bound PAP II, are the principal enzymes acting to metabolize TRH (649). TRH degradation in serum and many peripheral tissues is through the serum enzyme thyroliberinase (172). Each enzyme is described below in more detail.

Pyroglutamyl aminopeptidase I (PAP I) (EC 3.4.19.3) is a soluble cysteine protease that removes the N-terminal pyroGlu residue from TRH (649). It also cleaves peptides such as LHRH, NT, and bombesin. Prolyl endopeptidase (EC 3.4.21.26) is a soluble serine protease that cleaves on the carboxyl side of the TRH proline residue to generate TRH-OH (649). It also acts on other neuropeptides, including LHRH, NT, and SP. These two enzymes are present in many of the same tissues, so that both CHP and TRH-OH can be generated from TRH.

Displaying greater substrate specificity is the ectoenzyme pyroglutamyl aminopeptidase II (PAP II) (EC 3.4.19.6) (650). Like PAP I, this 260-kDa metalloenzyme removes pyro-Glu from TRH. The distinguishing features of PAP II are its greater substrate specificity and being membrane bound. PAP II is present in CNS synaptosomal fractions, in adeno-hypophyseal plasma membrane, and liver and serum particulate fractions. Highest activity is observed in the hippocampus and cerebral cortex. PAP II has been identified in many species, being highest in rabbit CNS, and in most cases

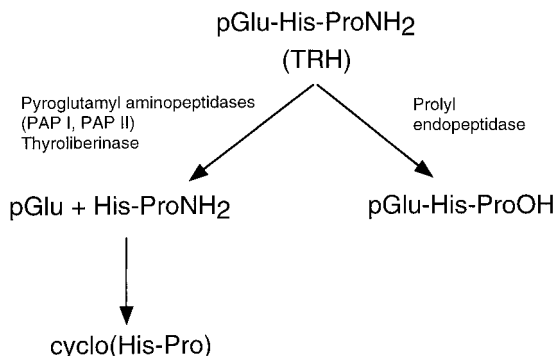


FIG. 10. Metabolic degradation of TRH by different TRH-degrading enzymes.

retains similar features. However, its substrate specificity in bovine synaptosomes, where it has been extensively characterized, is not as narrow as in other species (651). The localization of PAP II in the CNS is consistent with its proposed role in the degradation of synaptic TRH (652).

In the hypothalamus, PAP II activity is maximal at day 8 after birth, decreasing to adult values at day 45, while in the adenohypophysis it appears at day 8, peaks at day 30, and then decreases to adult values. In addition, thyroid hormone regulates the PAP II in the anterior pituitary but not in the brain (653, 654). In the PVN, TRH levels and PAP II activity do not correlate during pregnancy and lactation, indicating that PAP II is not the principal determinant of TRH levels (655). PAP II activity does vary with the estrous cycle (655). Furthermore, in brain areas other than the hypothalamus, PAP II activity decreases from days 9–20 coincident with increases in TRH and decreases in CHP (656), indicating PAP II activity can be a critical determinant of TRH steady state levels in some tissues. In sum, it appears that PAP II in areas under prominent endocrine control, such as the pituitary and PVN, subserves a different role than that in nonendocrine tissues.

Thyroliberinase, a fourth TRH-degrading enzyme, present in serum, is similar to PAP II but does not have the trans-membrane anchor of PAP II. Like PAP II, thyroliberinase displays greater substrate specificity than PAP I or prolyl endopeptidase (657, 658). Thyroliberinase may be regulated by thyroid hormone; TRH half-life ranges from about 2 min in the plasma of thyrotoxic animals to 6 min in hypothyroid animals. In humans, the half-life of TRH is similar (172).

In studies examining the ontogeny of TRH catabolizing enzymes in pancreas, PAP I and prolyl endopeptidase are detected at early stages of rat pancreatic development, while PAP II remains undetectable. PAP I-specific activity increases until day 3 and decreases after day 5, and prolyl endopeptidase levels peak at 20 days. Because this development does not parallel that seen for TRH levels, it appears that TRH levels in neonatal rat pancreas are principally determined by biosynthetic rates (659).

The physiological significance of the soluble enzymes PAP I and prolyl endopeptidase within the brain and spinal cord are unclear, since in the case of neurotransmitter inactivation, TRH is probably degraded outside the neuron by ectoenzymes located on the cell surface, or within lysosomes after endocytosis. Membrane-bound ectoenzymes that are specific for TRH, such as PAP II, are more logically located for hydrolysis of synaptically released peptides. Soluble enzymes are better situated to control degradation of TRH during its transport in the hypophyseal portal vessels and systemic TRH. The exact mechanisms that control the amount of TRH that ultimately reaches the pituitary remain to be elucidated.

VIII. Concluding Remarks

This review describes the latest knowledge on pro-TRH research by providing new insights into its biosynthesis and processing, the neuroendocrine regulation thereof, the biological actions of its products, and the signal transduction and catabolic pathways used by those products. We empha-

size the wide diversity of non-TRH pro-TRH-derived peptides that are now known, and the intracellular pathways they take to reach maturity. The recent information available on tissue-specific processing of pro-TRH in the brain adds a further dimension to our understanding of differential processing and its importance in generating biological diversity.

Understanding of the role of non-TRH pro-TRH-derived peptides represents an exciting new frontier in pro-TRH research. During biosynthesis, these sequences within the precursor may function as structural or targeting elements that guide the folding and sorting of pro-TRH and its larger intermediates so that subsequent processing and secretion are properly regulated. The unique anatomical distribution of the pro-TRH end products, as well as regulation of their levels by neuroendocrine or pharmacological manipulations, described in this review, argues that these peptides will have unique biological roles. Some of these roles, such as for prepro-TRH₁₆₀₋₁₆₉, will be within the HPT axis, while many others will be unrelated to traditional thyroid function. This review also gathers together an extensive array of data indicating that TRH can function far beyond the HPT axis and should command significant future effort as a focus to develop new therapeutics. These therapeutics, in the form of TRH analogs or nonprotein peptidomimetics, and perhaps using novel delivery systems, will advance our ability to develop TRH and non-TRH pro-TRH-derived peptide agonists and antagonists that can target pro-TRH-derived peptides functioning in specific tissues or brain loci. It is hoped that these new drugs might provide novel treatment approaches for some of today's most difficult health and societal issues, including drug abuse, depression, chronic pain disorders, and sequelae of CNS injury.

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