

# Calcitonin and calcitonin receptors

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## Summary

**Calcitonin (CT) is a polypeptide hormone with 32 aminoacids syntetized primarily by the thyroid.**

Several evidences support the existence of nonthyroidal CT like peptide. The CT gene transcript also encodes a distinct peptide known as calcitonin gene related peptide (CGRP) which is a potent vasodilator and responsible for the stimulation of the glomerular filtration rate. In addition, a 37 aminoacid peptide amylin has been originally isolated by pancreatic  $\beta$ -cells. Amylin is able to inhibit insulin secretion, glucose transport into the skeletal musculature and gluconeogenesis. It is also able to inhibit gastric emptying. In the kidney it is able to modulate Calcium ( $\text{Ca}^{2+}$ ) excretion and increases renin activity. Finally, high affinity amylin receptors have been identified in the brain of the rat. The calcitonin receptor (CTR) is a member of a sub-family of the seven-transmembrane domain G-protein coupled receptor super family that includes several peptides. Members of this family have a similar structure with other seven-membrane-spanning domain G-protein coupled receptors.

The genetic contribution to osteoporosis susceptibility is well documented and many studies demonstrated that genetic factors play important roles in the regulation of bone metabolism. Restriction Fragment Length Polymorphisms (RFLPs) for the CTR gene have been described in the literature with a positive association with the lumbar bone mineral density (BMD), femoral neck BMD and with a lower incidence of vertebral fractures.

**KEY WORDS:** calcitonin, calcitonin receptor, calcitonin receptor polymorphisms, osteoporosis.

## Calcitonin

The calcitonin (CT) is a polypeptide hormone first discovered by Coop and colleagues in the course of investigations to identify hormones that regulate serum levels of calcium (1). CT is able to decrease blood calcium levels by direct inhibition of mediated bone resorption and by enhancing calcium excretion by

the kidney (2, 3). Human CT is a single-chain peptide of 32-aminoacid residues. The molecular mass of the hormone is 3418 Da. There is a disulfide bridge connecting the cysteines at position 1 and 7 to form a 7 amino-acid ring structure at the amino terminus. The precursor of CT, pre-procalcitonin (Pre-ProCT) contains 141 aminoacids (4).

Several evidences support the existence of non-thyroidal CT-like peptide. Indeed, Davis et al. (5) demonstrated that prostate gland was able to produce a large amount of CT. In addition, CT is present in the central nervous system (6). The protein sequence for the CT has been determined in several species. It is highly conserved within the N-terminal loop region but demonstrates divergence in the rest of the sequence (7). The mature human CT originates from the calcitonin-I (CALC-I) gene on chromosome 11. The CT gene family consists of four known genes (CALC-I to CALC-IV) that contain nucleotide sequence homologies (8). CALC-I is the only gene that produces CT. However, both CALC-I and CALC-II genes produce the hormones CT gene-related peptide type I and II (CGRP-I and II). The CALC-III is able to produce both CT and CGRP and finally CALC-IV produces amylin (Fig. 1) (8).

The biosynthetic secretory pathway for CT involves a complex series of progressive modifications: after the biosynthesis and folding of ProCT, subsequent proteolytic processing occurs both within the Golgi apparatus (9) and later within the secretory granules. It is likely that the cleavage of ProCT and consequent immature CT could be due to the action of a prohormone convertase enzymes (PCs). During early posttranscriptional

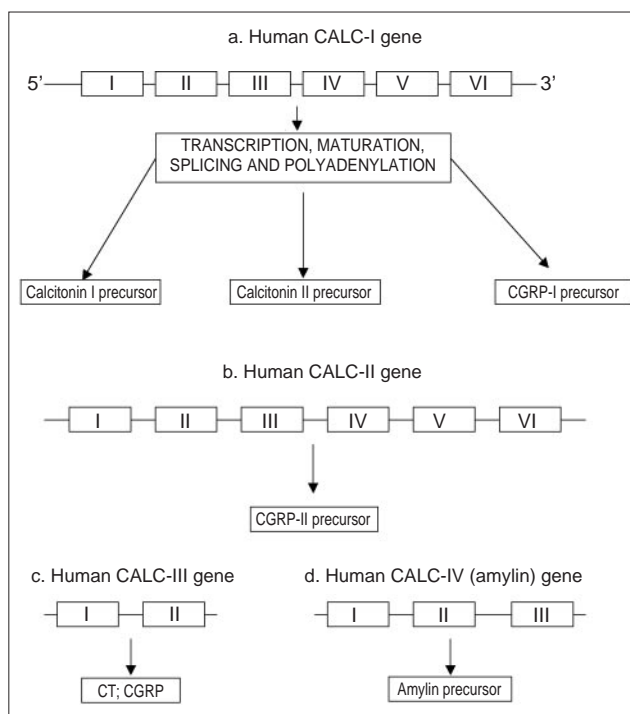


Figure 1 - The human calcitonin gene family.

processing, the nProCT segment may act as a signal for sorting its parent ProCT molecule to the nascent secretory vesicle of the regulatory pathway. Here, the prohormone undergoes further proteolytic cleavage to release CT (8). Chromogranin B may function as a helper protein to favor trans-Golgi sorting to the regulated secretory pathway (10). Within the newly formed secretion vesicles, the proteolytic cleavage releases immature CT. Then, as amidation proceeds, mature CT is produced and is progressively concentrated within the secretory vesicles (Fig. 2) (8, 11).

The biologically relevant effects of the CT appear in bone, kidney, central nervous system, respiratory tract and gastrointestinal and reproductive systems. Mature CT plays an important role in skeletal homeostasis, being a key modulator of bone resorption (8). It acts directly on osteoclast calcitonin receptor to inhibit bone resorption (1). The hormone inhibits bone resorption by inducing an acute quiescence of cell motility.

The physiological actions of CT are mediated primarily by the ability of its receptor to couple to at least two signal transduction pathways.

### Calcitonin-related peptide

CT is initially synthesized as a precursor containing 136 aminoacids and a leader sequence at its amino terminal region. The leader sequence is cleaved during transport of CT to the endoplasmic reticulum. The CT gene transcript also encodes a distinct peptide known as calcitonin gene-related peptide (CGRP). It is a potent vasodilator and responsible for the stimulation of the glomerular filtration rate (12). The tissue specificity of this transcript is regulated by alternative splicing of the CT gene (13). The pre-RNA of the CT/CGRP gene contain six exons and can be processed in two way with a regulatory event in the processing choice with inclusion or exclusion of the alternative 3' terminal exon 4. In particular, in parafollicular cells of the thyroid, 95% of the CT/CGRP pre-RNA is processed to include exon 4 followed by polyadenylation in order to produce CT. In the brain, 99% of the CT/CGRP pre-RNA is processed to exclude the exon 4 and include exon 5 and 6 with usage of the exon 6 polyadenylation site to produce CGRP (7, 13).

In addition, a 37 aminoacid peptide amylin has been originally

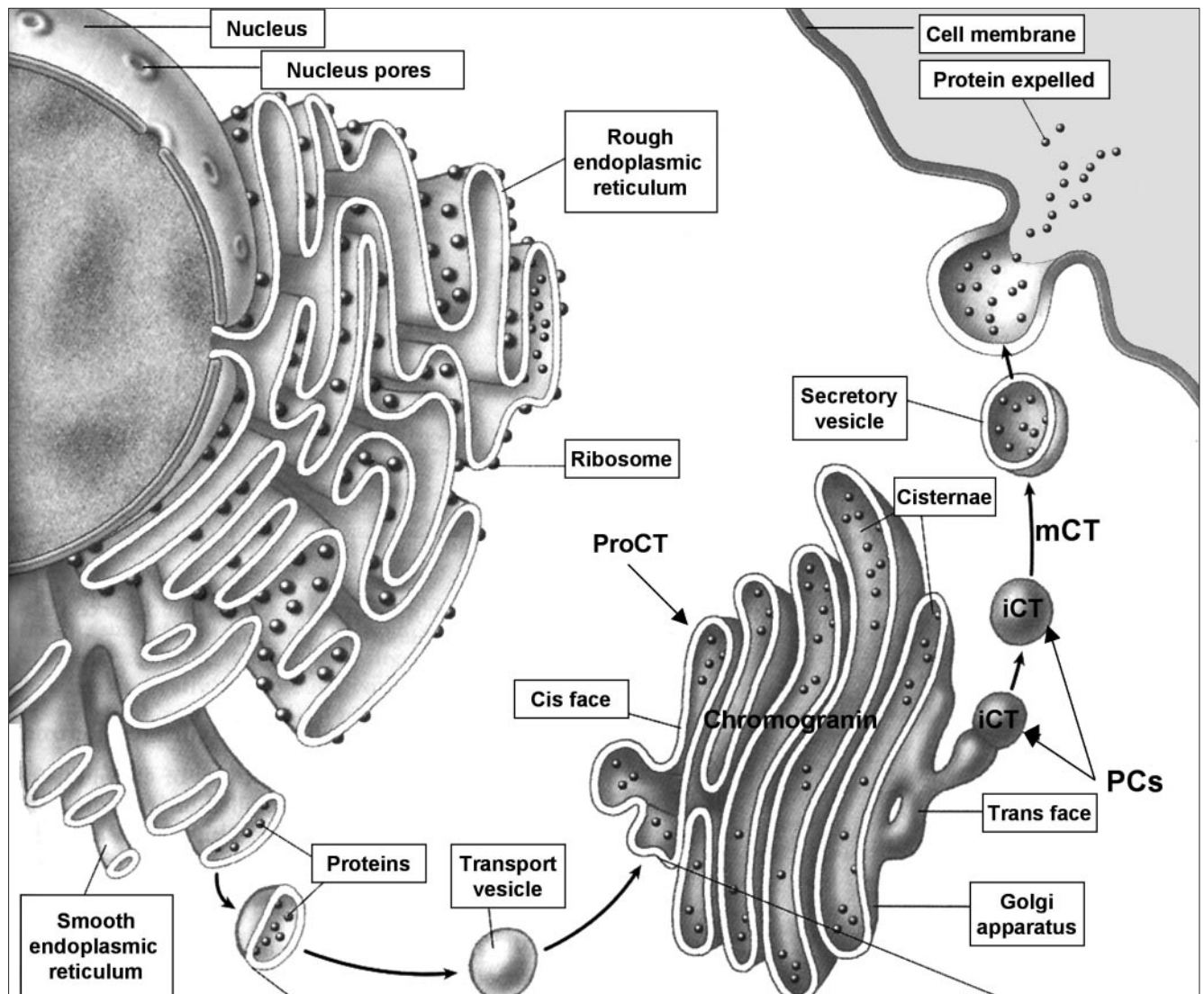


Figure 2 - The biosynthetic secretory pathway for CT. ProCT (ProCalcitonin) is cleaved into the Golgi apparatus by the action of a prohormone convertase enzymes (PCs). Consequently immature CT (iCT) present in the secretory vesicle, undergoes further proteolytic cleavage to release mature CT (mCT). Chromogranin B may function as a helper protein to favor trans-Golgi sorting to the regulated secretory pathway.

isolated by pancreatic  $\beta$ -cells (14). Calcitonin gene-related peptide (CGRP) and amylin are homologous 37 aminoacid peptides the gene for which probably has a common ancestral origin. Amylin is able to inhibit insulin secretion, glucose transport into the skeletal musculature and gluconeogenesis (7). In addition, amylin is able to inhibit gastric emptying. In the kidney it is able to modulate calcium ( $\text{Ca}^{2+}$ ) excretion and increases renin activity (15, 16). Finally, high affinity amylin receptors have been identified in the brain of the rat (17).

Amylin and CGRP have comparable effects. In skeletal muscle the primary effects are the inhibition of glycogen synthesis and stimulation of glycogenolysis with a reduction of glucose uptake and increased glycolysis. In the liver they stimulate the glucose output as a result of gluconeogenesis. By these ways amylin and CGRP act as noncompetitive antagonist of insulin and produce insulin resistance (19). In addition CGRP is potent vasodilator and its receptors are distributed throughout the vascular system. CGRP is also an abundant peptide in the nervous system suggesting a neuromodulatory role. Amylin and CGRP increase renin secretion. CGRP is also able to inhibit bone resorption. There is also evidence that CGRP may act on osteoclast precursors (20). Amylin has a similar effect but only at 100 fold higher concentrations. On the other hand, animal studies demonstrated that both amylin and CGRP are able to stimulate the osteoblast activity (18). However, what role amylin and CGRP play in normal bone metabolism and bone pathology remains to be determined. The effects of both amylin and CGRP are, broadly, to increase bone formation and reduce bone resorption. The most recently discovered member of the calcitonin peptide family is the adrenomedullin (AM). It has been isolated for the first time from a human phecomocytoma and it is expressed in the adrenal medulla (20). It is clear that either activation or disruption of AM signaling might contribute to many pathological conditions including hypertension, congestive heart failure, pulmonary hypertension, neoplastic growth, and inflammatory diseases (21).

### Calcitonin receptors

The activities of CT are mediated by high affinity calcitonin receptors (CTRs). The CTR is a member of a subfamily of the seven-transmembrane domain G-protein coupled receptor superfamily that includes several peptides. Members of this family have a similar structure with other seven-membrane-spanning domain G-protein coupled receptors. Lin et al. cloned the porcine CTR (pCTR) cDNA for the first time in 1991 (22). This receptor was characterized by a long NH<sub>2</sub>-terminal domain that was extracellular. It was similar to parathyroid/parathyroid hormone-related peptide receptor and the secretin receptor (7). Subsequent cloning of the pCTR gene demonstrated it is approximately 70 kb in length and contains at least 14 exons, 12 of which encode the protein. The human CTR was cloned from an ovarian carcinoma cell line (BIN-67) (23). Different isoforms of CTR resulting from alternative splicing of the gene have been described in various animal species with differential tissue expression transcripts (24) and different signaling properties (25). Two different isoforms have been described in human giant cell tumor of bone (26). The first isoform, designed as GC-10, differs from the previously described ovarian-human CTR gene (26) in the 5'-region in that it lacks a 71-bp segment, while being almost identical in the 3'-region. The second human giant cell tumor-CTR cDNA variant, indicated as GC-2, lacks 71-bp 5' insert but also 48 nucleotides encoding part of the first intracellular domain (26). It is likely that differential expression of CTR isoforms could be a mechanism of regulation of biological responses to CT. Shift in the predominant CTR isoform(s) could in part explain the variable responsiveness to

CT in patients with high turnover metabolic bone disease (26). The CTR gene has been mapped to chromosome 7q21.3 (27).

### Calcitonin and cell signalling

The principal mechanism of action of CT is due to the ability of its receptor to couple at least two signal transduction pathways. One of the most important pathways is coupled with the cAMP signal transduction. However, CTRs can also couple to the phospholipase C (PLC) enzyme pathway. The PLC pathway, as with the cAMP pathway, can be initiated by the coupling of receptors to multiple G-proteins. Activation of the PLC causes the release of  $\text{Ca}^{2+}$  from intracellular stores and promotes an influx of external calcium (7).

In addition, CTRs are able to activate the phospholipase D (PLD) (7). Finally, Chen et al. demonstrated that CT when bound to CTRs stimulates Shc tyrosine phosphorylation of MAPK Erk1/2 (28).

### Calcitonin activities

#### Bone

Osteoclasts are the major target for the action of CT. It is able to interfere with osteoclast differentiation from precursor cells and fusion of mononucleated precursors to form multinucleated cells in bone marrow cultures (29). CT plays an important role in skeletal homeostasis, being a key modulator on bone resorption (8). It acts directly on CTRs to inhibit bone resorption by inducing contraction and inhibits osteoclast motility (Q effect) occurring within 1 min and this is followed by a more gradual retraction of the osteoclasts (R effect) (7, 8, 30). Both cAMP and intracellular calcium ( $\text{Ca}^{2+}$ ) are second messengers for the Q and R effects, and both are G-protein mediated (8). CT inhibits also other components of the osteoclast, such as the release of acid phosphatase and the expression of carbonic anhydrase II (31, 32).

#### Kidney

The kidney is the principal site of CT degradation by neutral endopeptidase (NEP) (8). In normal men the effect of CT on the kidney is to stimulate diuresis and increases the fractional excretion rate of sodium and chloride. In addition, in urine a calcium and phosphate excretion increases. In hypercalcemic patients with metastatic bone disease, the administration of CT induces a rapid fall in serum calcium due primarily to inhibition of renal tubular reabsorption (7).

#### Central nervous system

CT has specific binding sites within the central nervous system. The intracerebral injection of CT suppresses food and water intake in rats (33). In the human, large doses of salmon CT reduce the serum concentrations of testosterone, LH and FSH probably acting via hypothalamic level (8, 34) and the chronic administration to humans with migraine headaches increases the levels of  $\beta$ -endorphin, as well as ACTH and cortisol (8).

#### Breast cancer cell lines

It is known that cell proliferation was inhibited when human breast cancer cells in culture were treated with CT (35, 36). Autoradiographic and radioligand-binding techniques with iodinated CT have identified high-affinity CTRs in the human breast cancer cell lines T47D and MCF-7 (36). In a recent study Wang et al. have demonstrated that CTR mRNA is constantly expressed in normal ductal epithelium and in breast cancer. In addition a decrease of CTR expression was found more often in cases with lymph node metastasis suggesting that CTR

might be of great potential significance in breast cancer progression (36).

#### Gastrointestinal system

In human, CT increases gastric acid and pepsin secretion and decreases pancreatic amylase and pancreatic polypeptide. At high concentrations, CT increases the net secretion of water and electrolytes from the human jejunum and ileum, and has been postulated that these effects may explain the presence of diarrhea seen in some patients affected by medullary thyroid carcinoma (8).

#### Reproductive system

Sperm contain a very high concentration of CT which is about 40 times that found in the serum. *In vitro* experiments showed that CT likely acts as an endogenous regulator of sperm fertilizing ability *in vivo* (7, 37). In addition, CT occurs in both the uterus and placenta. The intraplacental presence of CT receptors, suggest a role of CT in the regulation of placental function; however, small studies evaluated this possibility (8).

#### Respiratory system

9CT is found within the pulmonary neuroendocrine (PNE) cells, which are situated near the basement membrane and often extend to the lumen of the airway (8). CT affects transcellular and intracellular movements of calcium, and hence may exert and intrapulmonary paracrine action. In fact, CT is able to inhibit prostaglandin and thromboxane secretion (8) and increases the cartilaginous growth (8). Finally, CT blocks the bronchoconstrictor effects of bombesin-like peptides and substance P.

### Calcitonin receptor polymorphisms and osteoporosis

Osteoporosis is a common skeletal disease characterized by an excessively fragile skeleton and susceptibility to fracture (38). It is a multifactorial disease and several environmental and genetic factors play an important role. The genetic contribution to osteoporosis susceptibility is well documented and many studies demonstrated that genetic factors play important roles in determining population variation of Bone Mineral Density (BMD) (39), bone quality (40), and osteoporotic fractures (41, 42). The genes that regulate BMD and bone fragility are potentially important targets for the design of new drugs that can be used to treat bone disease. Genetic markers will also be used as diagnostic tools in the assessment of individuals at risk of developing osteoporotic fractures and could be used to help target preventive therapies to those individuals who are at risk of fracture. In addition, genetic profiling could be useful to distinguish treatment responders from non-responders and to identify patients who might be at risk of developing unwanted side effects.

One of the gene that can be involved in this process is the CTR gene. A Restriction Fragment Length Polymorphism (RFLP) for the CTR gene has been for the first time described in 1998 using Taq I endonuclease (43). The authors found a positive association between this polymorphism and lumbar spine bone mineral density (LS-BMD). However, no significant variation of BMD among the genotypes was observed in the femoral neck BMD (43). In addition, Nakamura et al. described a polymorphic site at the CTR gene by Alu I restriction enzyme at the 1377th nucleotide expressing either proline (CC genotype) or leucine (TT genotype) (44). They observed that in the Japanese population the most represented CTR was the CC genotype and the less represented was the TT genotype. Conversely in the Italian population the TC and TT allelic variants represented the most frequent genotypes, underlying the importance of ethnic origin as a variable that should be always taken in

consideration (45). In addition, Duncan's test used to compare the genotypes showed that TT genotype has significant lower lumbar BMD in comparison with CC genotype (Fig. 3). Taboulet et al. studied the distribution of these alleles in a cohort of 215 postmenopausal Caucasian women suffering or not from osteoporotic fractures. They found that Rr genotype (the TC genotype indicated by Masi et al. and Nakamura et al.) was associated with a higher femoral neck BMD in comparison with RR and rr genotypes. In addition Rr genotype was associated with a lower incidence of vertebral fractures (46). The authors emphasize the importance of the high conservation of the proline residue in all species studied with the exception of the human isoforms. The absence of the proline residue could alter the secondary structure of the calcitonin receptor, more so as two other proline residues border this region. Progressive truncation of the C-terminal tail of the porcine receptor complex reduced the magnitude of adenylate cyclase responses (47). This underlines the importance of the CTR C-terminal domain and suggests that proline/leucine mutation could alter receptor biological activity (46). In addition, heterozygotes could produce both alleles of the receptor, resulting in advantage as compared with homozygotes, especially as this genetic effect accounts for 13% of the total variance of BMD in covariance analysis (46). Recently, this polymorphism has been studied in a group of postmenopausal women in Taiwan (48) confirming the association between TT genotype and low bone mineral density at the lumbar spine and femoral neck. A lower lumbar spine BMD was also found in subjects with TT genotype affected by Juvenile Idiopathic Arthritis (49).

A complete polymorphism scan of the human calcitonin receptor gene identified 11 polymorphic sites in the gene and confirmed the presence of the previously identified nucleotide T1340C (codon 447) polymorphism. Because a leucine to proline change has the potential for significant structural alteration, receptor genes encoding either leucine or proline at residue 447 were transiently expressed in COS-7 cells to determinate the binding and functional consequences of this polymorphism. The presence of this SNP (single nucleotide polymorphism) appears to have no statistically significant difference with the receptor's ability to bind CT or signal when activated with the hormone (50). A study conducted in Italian men found that CC genotype was associated with lower BMD and with lower level of bone alkaline phosphatase and estradiol in comparison with TT genotype suggesting a depression of bone formation in these subjects (51). A correlation between CTR polymorphism and buccal margin bone loss has been observed in subjects undergoing to implants on maxillae and mandibles (52) and finally the CTR polymorphism has been indicated as an important genetic marker for severe periodontitis (53).

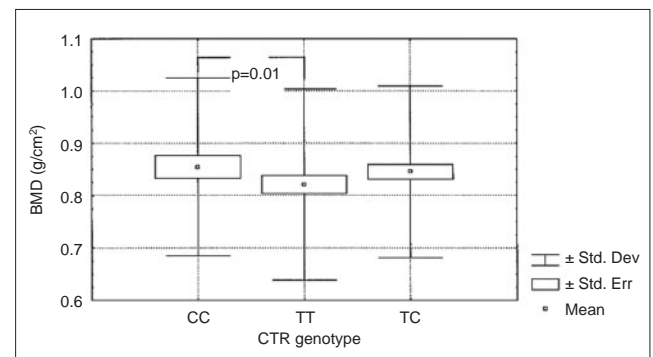


Figure 3 - Lumbar BMD values according to CTR genotypes. TT genotype had a significant lower lumbar BMD in comparison with CC genotype (Duncan's test  $p=0.01$ ).

Recently a functional role of CTR polymorphism has been observed by Zofkaova et al. (54). In particular, the authors postulated that the  $\beta$ 3-hydroxysteroid dehydrogenase activity could be associated with C allele at least in C19 steroids. However, these data were observed in a relatively small number of postmenopausal women and need confirm (54).

Research has recently begun to clarify the genes and genetic variants that predispose to osteoporosis and regulation of bone mass. CTR polymorphism together with other genes could be associated with the identification of genetic markers for assessment of fracture risk and the identification of novel molecular targets for the design of drugs that can be used to treat bone disease.

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