

# Gene Section

## Review

# MST1R (Macrophage stimulating 1 receptor)

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Published in Atlas Database: January 2001

Online updated version: <http://AtlasGeneticsOncology.org/Genes/RONID287.html>

DOI: 10.4267/2042/37697

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## Identity

**Other names:** C-MET-related tyrosine kinase (RON); RON protein tyrosine kinase (RON);; Macrophage stimulating protein receptor (MSP-receptor)

**HGNC (Hugo):** MST1R

**Location:** 3p21.31

**Local order:** Between LIMD1 and CCXCR1; between D3S1568 and D3S3822

## DNA/RNA

### Description

Twenty coding exons. All exons are small in size, ranging from 93 bp to 253 bp, with the exception of exon 1 (>1 kb). Exon 1, 2 and 3 code for the SEMA domain of the RON protein (red). Exons 4 codes for a PSI domain (orange), a modular structure about 50 amino acid long containing eight conserved Cys residues, putatively involved in protein-protein interactions. The sequence between exon 4 and 12 codes for four repeated modular structures called IPT (yellow); these domains are found in cell surface receptors such as MET and RON as well as

in intracellular transcription factors where they are involved in DNA binding. Part of exon 12 codes for the transmembrane domain, (pink). Exons 14 to 20 codes for the kinase domain (blue). Four-digit numbers refer to splice sites location, based on RON cDNA sequence.

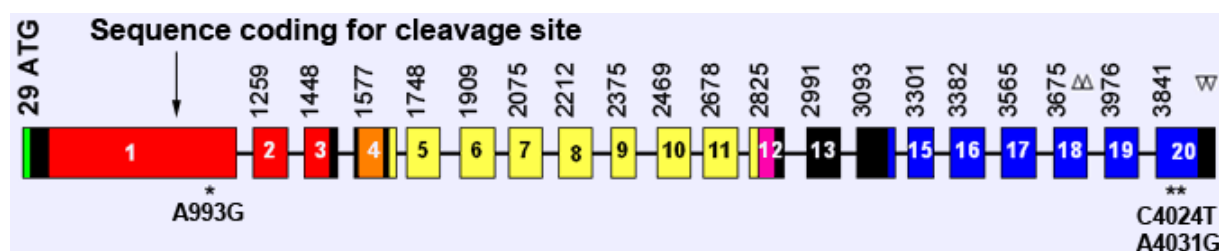
### Transcription

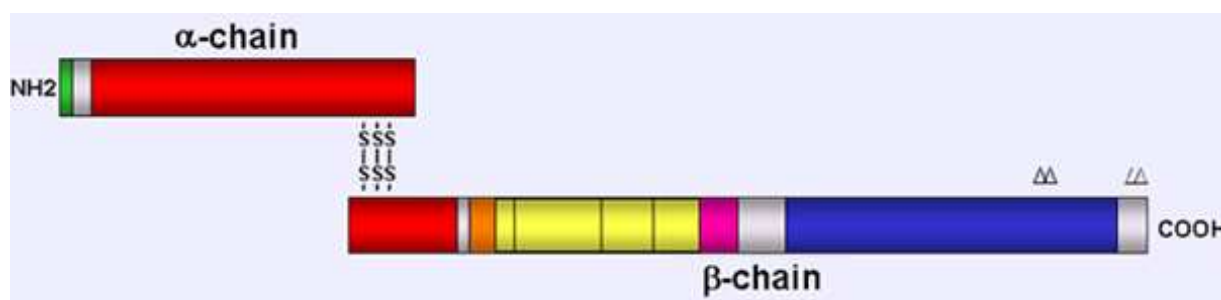
Two major transcripts are detected, respectively 4.5 kb and 2 kb. ORF: 4204 bp.

## Protein

### Description

The RON protein is a glycosylated heterodimeric protein composed of one  $\alpha$ - (35 kD) and one  $\beta$ -chain (150 kD) linked by an unknown number of disulfide bonds. The two chains derive from a single-chain precursor of about 185 kD that undergoes proteolytic cleavage at the basic amino acid site KRRRR. The  $\alpha$ -chain is extracellular. The  $\beta$ -chain has an extracellular part, a one-pass transmembrane helix and an intracellular part containing the tyrosine kinase domain. The first 24 amino acids made the putative signal peptide (green).





The SEMA domain (consisting of most of  $\alpha$ - and part of  $\beta$ - chain) contains the ligand (MSP) binding pocket (unpublished data). Tyrosine residues 1238 and 1239 (upward arrowheads in the figure) are essential for up-regulation of RON catalytic activity. Tyrosine residues 1353 and 1360 (downward arrowheads, in the figure) make a docking site that mediates high affinity interactions with multiple SH2-containing signal transducers.

### Expression

RON is expressed in human keratinocytes (it was initially cloned from a keratinocytes cDNA library). By Northern blot was found expressed in the following normal human tissues: skin, lung, bone marrow, small intestine, heart, pancreas, thyroid, prostate, testis (unpublished data), colonic mucosa and in a variety of cell types: granulocytes and monocytes, hematopoietic cells such as erythroid and myeloid progenitor cells, macrophages, osteoclasts, bone marrow megakaryocytes, epithelial and neuroendocrine cells.

### Localisation

Transmembrane protein.

### Function

The ligand for RON is MSP. Originally, MSP was described as a serum factor enhancing the chemotactic response of murine peritoneal macrophage to the C5a fraction of complement, but RON/MSP complex has a much broader spectrum of activity. Ligand-stimulated RON activates the pathways regulating cell adhesion and motility, growth and survival. STK (the mouse ortholog) is essential for peri-implantation development during gestation, as STK-deficient mice (STK<sup>-/-</sup>) are viable only through the blastocyst stage. Hemizygous mice (STK<sup>+/-</sup>) grow to adulthood; however, they are highly susceptible to endotoxic shock and appear to be compromised in their ability to down-regulate nitric oxide production. These results suggest STK has a limiting role not only in the inflammatory response but also in early mouse development.

### Homology

RON belongs to the MET receptor tyrosine kinase (RTK) family. On the basis of the presence of multiple PSI domains and a SEMA domain, it has been

proposed that plexins, MET RTK family and VESPR (virus-encoded semaphorin receptor) are classified as semaphorins. RON orthologs have been identified in mouse (STK), chicken (c-sea) and *Xenopus*.

## Mutations

### Germinal

Several Single Nucleotide Polymorphisms (SNPs) were found in healthy CEPH individuals: A993G:Gln322Arg (index of heterozygosity: 0.28); C4024T (same-sense variant, index of heterozygosity: 0.03); A4031G: Arg1344Gly (index of heterozygosity: 0.46).

### Somatic

T915C: Leu296Pro was found in the tumor DNA of one single patient affected with adenocarcinoma of the lung. The mutated protein is not constitutively activated. The mutation has no causative role in the disease. Experimental introduction in the RON kinase domain of amino acid substitutions D1232V and M1254T - initially found in the oncogenes KIT, RET and MET, involved respectively in mastocytosis, Multiple Endocrine Neoplasia type 2B and renal papillary carcinoma - results in activation of oncogenic capacity and triggers a strong metastatic activity of RON. Expression of these RON mutants causes cellular accumulation of  $\beta$ -catenin via inhibition of its association with the axin/GSK complex and subsequent protection from proteasomal degradation (Danilkovitch-Miagkova, personal communication).

## Implicated in

RON was found over-expressed in infiltrating breast carcinomas. A constitutively activated splicing variant of RON (lacking exon 11) was found in the gastric carcinoma cell line KATO-III. This variant induces activation of cell dissociation, motility and invasion of extracellular matrices. The same variant was found in malignant colonic mucosa. Another splicing variant, lacking exons 5 and 6, was found in the human colon carcinoma cell line HT-29. Truncated STK - the mouse RON ortholog - confers susceptibility to Friend virus-induced erythroleukemia in mice, and c-sea, the avian ortholog, causes erythroblastosis in chickens.

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*This article should be referenced as such:*

Angeloni D, Lerman MI. MST1R (Macrophage stimulating 1 receptor). *Atlas Genet Cytogenet Oncol Haematol.* 2001; 5(1):23-26.

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