Gene name: TREM1

External Ids for TREM1 Gene: HGNC: 17760 NCBI Gene: 54210 Ensembl: ENSG00000124731

OMIM®: 605085 UniProtKB/Swiss-Prot: Q9NP99

NCBI Gene Summary: This gene encodes a receptor belonging to the lg superfamily that is expressed on myeloid cells. This protein amplifies neutrophil and monocyte-mediated inflammatory responses triggered by bacterial and fungal infections by stimulating release of pro-inflammatory chemokines and cytokines, as well as increased surface expression of cell activation markers. Alternatively spliced transcript variants encoding different isoforms have been noted for this gene.

GeneCards Summary: TREM1 (Triggering Receptor Expressed On Myeloid Cells 1) is a Protein Coding gene. Diseases associated with TREM1 include Maxillary Sinusitis and Pneumonia. Among its related pathways are Innate Immune System and Toll-like receptor signaling pathway. Gene Ontology (GO) annotations related to this gene include *signaling receptor activity* and *scaffold protein binding*.

UniProtKB/Swiss-Prot Summary: [Isoform 1]: Cell surface receptor that plays important roles in innate and adaptive immunity by amplifying inflammatory responses (PubMed:10799849, 21393102). Upon activation by various ligands such as PGLYRP1, HMGB1 or HSP70, multimerizes and forms a complex with transmembrane adapter TYROBP/DAP12 (PubMed:17568691, 25595774, 29568119). In turn, initiates a SYK-mediated cascade of tyrosine phosphorylation, activating multiple downstream mediators such as BTK, MAPK3 or phospholipase C-gamma (PubMed:14656437, 21659545). This cascade promotes the neutrophil- and macrophage-mediated release of pro-inflammatory cytokines and/or chemokines, as well as their migration and thereby amplifies inflammatory responses that are triggered by bacterial and fungal infections (PubMed:17098818, 17568691). By also promoting the amplification of inflammatory signals that are initially triggered by Toll-like receptor (TLR) and NOD-like receptor engagement, plays a major role in the pathophysiology of acute and chronic inflammatory diseases of different etiologies including septic shock and atherosclerosis (PubMed:11323674, 21393102). (TREM1_HUMAN,Q9NP99) [Isoform 2]: Acts as a decoy receptor, counterbalancing TREM1 pro-inflammatory activity through the neutralization of its ligand. (TREM1_HUMAN,Q9NP99).

Gene Name: TREM1 (Triggering Receptor Expressed on Myeloid Cells 1).

TREM1 is an immunoreceptor primarily expressed on myeloid cells, such as: Neutrophils,

Monocytes/macrophages. It acts as an amplifier of the innate immune response, especially during bacterial or fungal infections.

Molecular Function

- TREM1 is part of the immunoglobulin superfamily.
- It does not have a signaling domain itself, but signals via DAP12 (TYROBP), a protein with an ITAM (immunoreceptor tyrosine-based activation motif) that triggers intracellular activation.
- When activated, it induces secretion of pro-inflammatory cytokines like: TNF-α, IL-6, IL-8, MCP-1.

TREM1 in Sepsis

TREM1 is strongly upregulated during sepsis and other systemic inflammatory responses, particularly in:

- Sepsis-induced acute lung injury, Bacteremia and Septic shock. It works synergistically with Toll-like receptors (TLRs) — especially TLR4, amplifying their response to pathogen-associated molecular patterns (PAMPs).
- Inhibiting TREM1 signaling (such as, with TREM1 decoy peptides like LP17) in mouse models:
 - Reduces cytokine storm
 - Improves survival
 - Prevents tissue damage

Diagnostic/Prognostic Value

- sTREM1 (soluble form of TREM1) can be measured in plasma or serum, and:
 - Is elevated in sepsis.
 - Correlates with severity (SOFA score, mortality risk).
 - o May help differentiate bacterial infection from non-infectious inflammation.

Clinical trials are exploring sTREM1 as a biomarker for: Early sepsis detection, Monitoring disease progression, Assessing response to treatment

Supporting Literature Doi: 10.1016/j.cellimm.2011.10.006

Doi: 10.3389/fimmu.2022.907387 Doi:10.1002/eji.200636387

Enrichr-KG:

