

Recombinant Rat RAGE Protein

For Research Use Only

1. Basic Product Information

Catalog No.: REP52500

Size: 10ug, 100ug

Protein Name: Advanced glycation end product-specific receptor (Receptor for advanced glycation end products)

Synonym: Ager, Rage

UniProt ID: Q63495

UniProt Link: www.uniprot.org/uniprotkb/Q63495/entry

Species Source: Rat

Expression System: HEK293T eukaryotic cell expression system Sequence

Length: 402 AA

Molecular Weight: 43 KD

Protein Tag: C-terminal His6

2. Amino Acid Sequence

MPTGTVARAWVLVLALWGAVAGGQNITARIGEPLMLSCKGAPKKPTQKLEWKLNTGRTEAWKVLSP
QGDWPDSVARILPNGSLLLPAIGIVDEGTFRCRATNRLGKEVKSNYRVRVYQIPGKPEIVNPASELTA
NVPNKVGTCTVSEGSYPAGTLSWHLDGKPLIPDGKGTVVKEETRRHPETGLFTLRSELTVTTPAQGGT
TPTYSCSFSGLPRRRPLNTAPIQPRVREPLPPEGIQLLVEPEGGTVAPGGTVTLTCAISAQPPPQIH
WIKDGTPLPLAPSPVLLLPEVGHEDEGIYSCVATHPSHGPQESPPVNIRVTETGDEGQAAGSVDGSG
LGTALALGILGGLGIAALLIGAILWRKRQPRLEERKAPESQEDEEERAELNQSEEAEMPENGAGGP

3. Buffer Composition

Liquid format: PBS-based buffer containing 50% glycerol. Glycerol concentration can be customized upon customer request; please state special requirements when placing your order.

Lyophilized format: Buffer before lyophilization is PBS supplemented with 6% trehalose.

4. Storage Conditions



Upon receipt, store the product at -20 °C or -80 °C. Aliquot the product for on-demand use to preserve protein activity and prevent denaturation and inactivation caused by repeated freeze-thaw cycles.

5. Product Description

This product is recombinant Advanced glycation end product-specific receptor (Receptor for advanced glycation end products) produced using the HEK293T eukaryotic cell expression system. It is suitable for in-vitro functional assays, protein-protein interaction studies, antibody preparation and other biomedical research applications.

Protein Function: Cell surface pattern recognition receptor that senses endogenous stress signals with a broad ligand repertoire including advanced glycation end products, S100 proteins, high-mobility group box 1 protein/HMGB1, amyloid beta/APP oligomers, nucleic acids, histones, phospholipids and glycosaminoglycans (PubMed:19910580, PubMed:28627626). Advanced glycation end products (AGEs) are nonenzymatically glycosylated proteins which accumulate in vascular tissue in aging and at an accelerated rate in diabetes. These ligands accumulate at inflammatory sites during the pathogenesis of various diseases including diabetes, vascular complications, neurodegenerative disorders and cancers, and RAGE transduces their binding into pro-inflammatory responses. Upon ligand binding, uses TIRAP and MYD88 as adapters to transduce the signal ultimately leading to the induction of inflammatory cytokines IL6, IL8 and TNFalpha through activation of NF-kappa-B. Interaction with S100A12 on endothelium, mononuclear phagocytes, and lymphocytes triggers cellular activation, with generation of key pro-inflammatory mediators (By similarity). Interaction with S100B after myocardial infarction may play a role in myocyte apoptosis by activating ERK1/2 and p53/TP53 signaling (By similarity). Contributes to the translocation of amyloid-beta peptide (ABPP) across the cell membrane from the extracellular to the intracellular space in cortical neurons. ABPP-initiated RAGE signaling, especially stimulation of p38 mitogen-activated protein kinase (MAPK), has the capacity to drive a transport system delivering ABPP as a complex with RAGE to the intraneuronal space. Participates in endothelial albumin transcytosis together with HMGB1 through the RAGE/SRC/Caveolin-1 pathway, leading to endothelial hyperpermeability. Mediates the loading of HMGB1 in extracellular vesicles (EVs) that shuttle HMGB1 to hepatocytes by transferrin-mediated endocytosis and subsequently promote hepatocyte pyroptosis by activating the NLRP3 inflammasome. Binds to DNA and promotes extracellular hypomethylated DNA (CpG DNA) uptake by cells via the endosomal route to activate inflammatory responses (By similarity). Mediates phagocytosis by non-professional phagocytes (NPP) and this is enhanced by binding to ligands including RNA, DNA, HMGB1 and histones (By similarity). Promotes NPP-mediated phagocytosis of *Saccharomyces cerevisiae* spores by binding to RNA attached to the spore wall (By similarity). Also promotes NPP-mediated phagocytosis of apoptotic cells (By similarity). Following DNA damage, recruited to DNA double-strand break sites where it colocalizes with the MRN repair complex via interaction with double-strand break repair protein MRE11 (By similarity). Enhances the endonuclease activity of MRE11, promoting the end resection of damaged DNA (By similarity). Promotes DNA damage repair in trophoblasts which enhances trophoblast invasion and contributes to placental development and maintenance (By similarity). Protects cells from DNA replication stress by localizing to damaged replication forks where it stabilizes the MCM2-7 complex and promotes faithful progression of the replication fork (By similarity).

Subcellular Localization: Cell membrane ; Single-pass type I membrane protein . Cell projection, phagocytic cup . Early endosome . Nucleus . Note=Nuclear translocation is enhanced by irradiation, hypoxia and reperfusion injury to brain or kidney.

6. Notes

- Avoid repeated freeze-thaw cycles to maintain protein stability and biological activity.
- Please specify special buffer-composition requirements at the time of order placement.



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- For research use only. Not intended for clinical diagnosis, therapeutic use or commercial manufacturing.