

DESCRIPTION

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|---------------|--|---------|---|
| Source | Human embryonic kidney cell, HEK293-derived | | |
| | Human HGF R α (Glu25-Arg307) Accession # P08581 | | |
| | Human HGF R β (Ser308-Thr932) Accession # P08581 | HIEGRMD | Human IgG ₁ (Pro100-Lys330) |
| | N-terminus | | C-terminus |

N-terminal Sequence Analysis Glu25 & Ser308

Structure / Form Tetramer containing two disulfide-linked α and β subunits

Predicted Molecular Mass 33 kDa & 96 kDa

SPECIFICATIONS

| | |
|------------------------|---|
| SDS-PAGE | 34-48 kDa and 103-126 kDa, reducing conditions |
| Activity | Measured by its ability to bind immobilized recombinant human HGF in a functional ELISA with an estimated K_d <0.15 nM. |
| Endotoxin Level | <0.10 EU per 1 μ g of the protein by the LAL method. |
| Purity | >85%, by SDS-PAGE with silver staining. |
| Formulation | Lyophilized from a 0.2 μ m filtered solution in PBS. See Certificate of Analysis for details. |

PREPARATION AND STORAGE

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|--------------------------------|---|
| Reconstitution | Reconstitute at 250 μ g/mL in PBS. |
| Shipping | The product is shipped at ambient temperature. Upon receipt, store it immediately at the temperature recommended below. |
| Stability & Storage | <p>Use a manual defrost freezer and avoid repeated freeze-thaw cycles.</p> <ul style="list-style-type: none"> ● 12 months from date of receipt, -20 to -70 °C as supplied. ● 1 month, 2 to 8 °C under sterile conditions after reconstitution. ● 3 months, -20 to -70 °C under sterile conditions after reconstitution. |

BACKGROUND

HGF R, also known as Met (from *N*-methyl-*N*-nitro-*N*-nitrosoguanidine induced), is a glycosylated receptor tyrosine kinase that plays a central role in epithelial morphogenesis and cancer development. HGF R is synthesized as a single chain precursor which undergoes cotranslational proteolytic cleavage. This generates a mature HGF R that is a disulfide-linked dimer composed of a 50 kDa extracellular α chain and a 145 kDa transmembrane β chain (1, 2). The extracellular domain (ECD) contains a seven bladed β -propeller sema domain, a cysteine-rich PSI/MRS, and four Ig-like E-set domains, while the cytoplasmic region includes the tyrosine kinase domain (3, 4). Proteolysis and alternative splicing generate additional forms of human HGF R which either lack of the kinase domain, consist of secreted extracellular domains, or are deficient in proteolytic separation of the α and β chains (5-7). The sema domain, which is formed by both the α and β chains of HGF R, mediates both ligand binding and receptor dimerization (3, 7). Ligand-induced tyrosine phosphorylation in the cytoplasmic region activates the kinase domain and provides docking sites for multiple SH2-containing molecules (8, 9). HGF stimulation induces HGF R down-regulation via internalization and proteasome-dependent degradation (10). In the absence of ligand, HGF R forms noncovalent complexes with a variety of membrane proteins including CD44v6, CD151, EGF R, Fas, Integrin $\alpha 6/\beta 4$, Plexins B1, 2, 3, and MSP R/Ron (11-18). Ligation of one complex component triggers activation of the other, followed by cooperative signaling effects (11 - 18). Formation of some of these heteromeric complexes is a requirement for epithelial cell morphogenesis and tumor cell invasion (11, 15, 16). Paracrine induction of epithelial cell scattering and branching tubulogenesis results from the stimulation of HGF R on undifferentiated epithelium by HGF released from neighboring mesenchymal cells (19). Genetic polymorphisms, chromosomal translocation, over-expression, and additional splicing and proteolytic cleavage of HGF R have been described in a wide range of cancers (1). Within the ECD, human HGF R shares 86%-88% aa sequence identity with canine, mouse, and rat HGF R.

References:

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