

## 32-6852: MMP14 Human

### Alternative Name :

Matrix Metalloproteinase 14, Matrix Metalloproteinase 14 (Membrane-Inserted), Membrane-Type-1 Matrix Metalloproteinase, Membrane Type 1 Metalloprotease, EC 3.4.24.80, MT-MMP 1, MT1-MMP, MMP-14, MMP-X1, MT1MMP, MTMMP1, Matrix Metalloproteinase 14 (Membrane-Inserted), Membrane-Type Matrix Metalloproteinase 1, Matrix Metalloproteinase-14, EC 3.4.24, MT-MMP, WNCHRS, Matrix metalloproteinase-14, Membrane-type matrix metalloproteinase 1.

### Description

Source: Escherichia Coli.

Sterile filtered colorless solution.

Matrix metalloproteinase-14 (MMP14), is a membrane-anchored zinc-binding endopeptidase which is expressed at the leading edge of different invasive carcinomas and also promotes tumor cell invasion through degradation of the extracellular matrix. MMP14 takes a vital part in extracellular matrix, ECM, remodeling by having the capability to degrade type I collagen, activate pro-MMP-2 and process cell adhesion molecules for instance CD44 and integrin alpha V. MMP14 is a key enzyme in many physiological as well as pathological processes for example angiogenesis & tumor invasion.

Matrix Metalloproteinase-14 Human Recombinant produced in E.Coli is a single, non- glycosylated polypeptide chain containing 264 amino acids and having a molecular mass of 29.6kDa. MMP14 is purified by proprietary chromatographic techniques.

### Product Info

<b>Amount :</b>	50 µg / 100 µg
<b>Purification :</b>	Greater than 95.0% as determined by:(a) Analysis by RP-HPLC.(b) Analysis by SDS-PAGE.
<b>Content :</b>	MMP14 is supplied as a 0.2 µm filtered solution containing 20mM Tris-HCl, pH 7.4, 30 % glycerol, 300mM NaCl, 3mM CaCl <sub>2</sub> and 10µM ZnCl <sub>2</sub> .
<b>Storage condition :</b>	Store at 4°C if entire vial will be used within 2-4 weeks.Store, frozen at -20°C for longer periods of time.For long term storage it is recommended to add a carrier protein (0.1% HSA or BSA).Avoid multiple freeze-thaw cycles.
<b>Amino Acid :</b>	ALASLGSAQS SSFSPEAWLQ QYGYLPPGDL RTHTQRSPQS LSAAIAMQK FYGLQVTGKA DADTMKAMRR PRCGVDPKFG AEIKANVRRK RYAIQGLKWQ HNEITFCIQN YTPKVGGEYAT YEAIRKAFRV WESATPLRFR EVPYAYIREG HEKQADIMIF FAEGFHGDST PFDGEGGFLA HAYFPGPNIG GDTHFDSAEP WTVRNEDLNG NDIFLVAVHE LGHALGLEHS SDPSAIMAPF YQWMDTENFV LPDDDRRGIQ QLYG.