## 32-2459: HTRA2 Recombinant Protein

Alternative Name Serine protease HTRA2 mitochondrial,EC 3.4.21.108, High temperature requirement protein A2,HtrA2,Omi : stress-regulated endoprotease,Serine proteinase OMI,Serine protease 25,OMI,PARK13,PRSS25.

## Description

Source : Escherichia Coli. HtrA2 Human Recombinant amino acids $134-458$ His-Tag fusion protein produced in E.Coli is a single, non-glycosylated polypeptide chain having a molecular mass of 36 kDa . The HtrA2 is purified by proprietary chromatographic techniques. HtrA2 also called Omi is a mammalian serine protease at high temperatures and has a chaperone activity at low temperature. The full-length HtrA2 is synthesized as a precursor protein and then targeted to the mitochondria where it is matured by the removal of N -terminal 133 residues. Mature HtrA2 consists of a putative transmembrane domain; an inhibitor of apoptosis protein (IAP)-binding motif; a single C-terminal PDZ domain that mediates protein-protein interactions. Recently, HtrA2 has known to contribute both to caspase-dependent and caspase-independent cell death.

## Product Info

| Amount : | $50 \mu \mathrm{~g}$ |
| :---: | :---: |
| Purification : | Greater than 95.0\% as determined by SDS-PAGE. |
| Content : | The protein ( $0.5 \mathrm{mg} / \mathrm{ml}$ ) contains 20 mM Tris- HCl buffer ( pH 8.0 ), $50 \mathrm{mM} \mathrm{NaCl}, 1 \mathrm{mM}$ DTT, and $20 \%$ Glycerol. |
| Storage condition : | Store at $4^{\circ} \mathrm{C}$ if entire vial will be used within $2-4$ weeks. Store, frozen at $-20^{\circ} \mathrm{C}$ 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. |
| Amino Acid : | MAVPSPPPAS PPSQYNFIAD VVEKTAPAVV YIEILDRHPF LGREVPISNG SGFVVAADGL |
|  | IVTNAHVVAD RRRVRVRLLS GDTYEAVVTA VDPVADIATL RIQTKEPLPT LPLGRSADVR |
|  | QGEFVVAMGS PFALQNTITS GIVSSAQRPA RDLGLPQTNV EYIQTDAAID FGNAGGPLVN |
|  | LDGEVIGVNT MKVTAGISFA IPSDRLREFL HRGEKKNSSS GISGSQRRYI GVMMLTLSPS |
|  | ILAELQLREP SFPDVQHGVL IHKVILGSPA HRAGLRPGDV ILAIGEQMVQ NAEDVYEAVR |
|  | TQSQLAVQIR RGRETLTLYV TPEVTEGSHH HHHH. |



