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## 32-3057: MAPKAPK3 Recombinant Protein

**Alternative** Name:

3PK,MAPKAP-K3,MAPKAP3,MAPKAPK-3,MK-3,MAP kinase-activated protein kinase 3,MAPK-activated

protein kinase 3,MAPKAP kinase 3,MAPKAPK-3,MK-3,MAPKAPK3.

## **Description**

Source: E.coli. MAPKAPK3 Human Recombinant produced in E.Coli is a single, non-glycosylated polypeptide chain containing 405 amino acids (1-382 a.a) and having a molecular mass of 45.4kDa. MAPKAPK3 is fused to a 23 amino acid His-tag at Nterminus & purified by proprietary chromatographic techniques. MAP kinase-activated protein kinase 3 (MAPKAPK3) is involved in inflammatory Reaction by regulating tumor necrosis factor (TNF) and IL6 production post-transcriptionally. MAPKAPK3 phosphorylates AU-rich elements (AREs)-binding proteins, like TTP/ZFP36, leading to control of stability and translation of TNF and IL6 mRNAs. Phosphorylation of TTP/ZFP36 (a major post-transcriptional regulator of TNF), promotes its binding to 14-3-3 proteins and reduces its ARE mRNA affinity resulting in inhibition of dependent degradation of ARE-containing transcript. MAPKAPK3 is activated by growth inducers and stress stimulation of cells.

## **Product Info**

Amount: 20 µg

Greater than 95% as determined by SDS-PAGE. Purification:

MAPKAPK3 protein solution (1mg/ml) containing 20mM Tris-HCl buffer (pH 8.0), 0.2M NaCl, 20% Content:

glycerol and 1mM DTT.

Store at 4°C if entire vial will be used within 2-4 weeks. Store, frozen at -20°C for longer periods of Storage condition:

time. For long term storage it is recommended to add a carrier protein (0.1% HSA or BSA). Avoid

multiple freeze-thaw cycles.

MGSSHHHHHH SSGLVPRGSH MGSMDGETAE EQGGPVPPPV APGGPGLGGA PGGRREPKKY **Amino Acid:** 

> AVTDDYQLSK QVLGLGVNGK VLECFHRRTG QKCALKLLYD SPKARQEVDH HWQASGGPHI VCILDVYENM HHGKRCLLII MECMEGGELF SRIQERGDQA FTEREAAEIM RDIGTAIQFL HSHNIAHRDV KPENLLYTSK EKDAVLKLTD FGFAKETTQN ALQTPCYTPY YVAPEVLGPE KYDKSCDMWS LGVIMYILLC GFPPFYSNTG QAISPGMKRR IRLGQYGFPN PEWSEVSEDA KQLIRLLLKT DPTERLTITQ FMNHPWINQS MVVPQTPLHT ARVLQEDKDH WDEVKEEMTS

ALATMRVDYD QVKIKDLKTS NNRLLNKRRK KQAGSSSASQ GCNNQ.

