

## 32-3048: MAPK1 His Recombinant Protein

**Alternative Name :** Mitogen-activated protein kinase 1, EC 2.7.11.24, Extracellular signal-regulated kinase 2, ERK-2, Mitogen-activated protein kinase 2, MAP kinase 2, MAPK 2, p42-MAPK, ERK1, ERK, p38, p40, p41, ERK2, MAPK2, PRKM1, PRKM2, P42MAPK, p41mapk.

### Description

Source : Escherichia Coli. MAPK1 Human Recombinant produced in E.Coli is a single, non-glycosylated, polypeptide chain containing 380 amino acids (1-360 a.a.) and having a molecular mass of 43.5 kDa. The MAPK1 is purified by proprietary chromatographic techniques. Mitogen-activated protein kinase 1 (MAPK1) is also known as 'extracellular signal-regulated kinase 2' (ERK2). Two similar (85% sequence identity) protein kinases were originally called ERK1 and ERK2. They were found during a search for protein kinases that are rapidly phosphorylated after activation of cell surface tyrosine kinases such as the epidermal growth factor receptor. Phosphorylation of ERKs leads to the activation of their kinase activity. The molecular events linking cell surface receptors to activation of ERKs are complex. It was found that Ras GTP-binding proteins are involved in the activation of ERKs. Another protein kinase, Raf-1, was shown to phosphorylate a 'MAPK kinase', thus qualifying as a 'MAPK kinase kinase'.

### Product Info

**Amount :** 25 µg  
**Purification :** Greater than 95.0% as determined by SDS-PAGE.  
**Content :** The MAPK1 solution contains 20 mM Tris-HCl buffer (pH 8.0), 1mM DTT and 10% glycerol.  
**Storage condition :** MAPK1 although stable 4°C for 4 weeks, should be stored desiccated below -18°C. For long term storage it is recommended to add a carrier protein (0.1% HSA or BSA). Please prevent freeze-thaw cycles.  
**Amino Acid :** MGSSHHHHHH SSGLVPRGSH MAAAAAAGAG PEMVRGQVFD VGPRYTNSY IGEGAYGMVC  
 SAYDNVNKVR VAIKKISPFE HQTQCRTL R EIKILLRFRH ENIGINDII RAPTIEQMKD VYIVQDLMET  
 DLYKLLKTQH LSNDHICYFL YQILRGLKYI HSANVLHRDL KPSNLLLNTT CDLKICDFGL ARVADPDHHDH  
 TGFLTEYVAT RWYRAPEIML NSKGYSID IWSVGCILAE MLSNRPIFPG KHYLDQLNHI LGILGSPSQE  
 MGSSHHHHHH SSGLVPRGSH MAAAAAAGAG PEMVRGQVFD VGPRYTNSY IGEGAYGMVC  
 SAYDNVNKVR VAIKKISPFE HQTQCRTL R EIKILLRFRH ENIGINDII RAPTIEQMKD VYIVQDLMET  
 DLYKLLKTQH LSNDHICYFL YQILRGLKYI HSANVLHRDL KPSNLLLNTT CDLKICDFGL ARVADPDHHDH  
 TGFLTEYVAT RWYRAPEIML NSKGYSID IWSVGCILAE MLSNRPIFPG KHYLDQLNHI LGILGSPSQE  
 DLNCIINLKA RNYLLSLPHK NKVPWNRLFP NADSKALDLL DKMLTFNPHK RIEVEQALAH PYLEQYYDPS  
 DEPIAEAPFK FDMELDDLPK EKLKELIFEE TARFQPGYRS.

