

P90775Hu02
Hexokinase 1 (HK1)
Organism: Homo sapiens (Human)
Instruction manual

FOR IN VITRO USE AND RESEARCH USE ONLY
NOT FOR USE IN DIAGNOSTIC OR THERAPEUTIC PROCEDURES

5th Edition (Revised in January, 2013)

[DESCRIPTION]

Protein Names: Hexokinase 1

Synonyms: HK1

Species: Human

Size: 100µg

Source: *Escherichia coli*-derived

Subcellular Location: Mitochondrion outer membrane.

[PROPERTIES]

Residues: Asp413~Val540 (Accession # P19367), with two N-terminal Tags, His-tag and GST-tag.

Grade & Purity: >95%, 43kDa as determined by SDS-PAGE reducing conditions.

Formulation: Supplied as lyophilized form in PBS, pH 7.4, containing 5% sucrose.

Endotoxin Level: <1.0 EU per 1µg (determined by the LAL method).

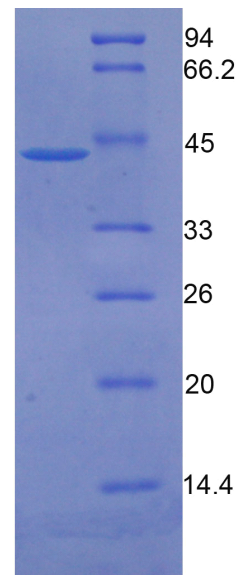
Applications: SDS-PAGE; WB; ELISA; IP.

(May be suitable for use in other assays to be determined by the end user.)




Predicted Molecular Mass: 41.7kDa

Predicted isoelectric point: 8.5

Human HK1 kDa



15% SDS-PAGE

Unique product Superb quality Client favorite Nicest service  ISO9001:2008;  ISO13485:2003; 

[PREPARATION]

Reconstitute in sterile PBS, pH7.2-pH7.4.

[STORAGE AND STABILITY]

Storage: Avoid repeated freeze/thaw cycles.

Store at 2-8°C for one month.

Aliquot and store at -80°C for 12 months.

Stability Test: The thermal stability is described by the loss rate of the target protein. The loss rate was determined by accelerated thermal degradation test, that is, incubate the protein at 37°C for 48h, and no obvious degradation and precipitation were observed. (Referring from China Biological Products Standard, which was calculated by the Arrhenius equation.) The loss of this protein is less than 5% within the expiration date under appropriate storage condition.

[SEQUENCES]

The target protein is fused with two N-terminal Tags, His-tag and GST-tag, its sequence is listed below.

MRNKKFELGLEFPNLPYYIDGDVKLTSMAIIRYIADKHNMLGGCPKERAEISMLEGAVLDIRYG
VSRIAYSKDFETLKVDFLSKLPEMLKMFEDRLCHKTYLNGDHVTHPDFMLYDALDVVLYMDPM
CLDAFPKLVCFKKRIEAIQIDKYLKSSKYIAWPLQGWQATFGGGDHPKSDGSTSGSGHHHH
HHSAGLVPRGSTAIGMKETAALKFERQHMDSPDLGTLEVLVFGPLGSEF-DGSLYKTH
PQYSRRFHKT LRRVLPDSV RFLSESGSG KGAAMVTAVA YRLAEQHRQI EETLAHFHLL
KDMILLEVKKR MRAEMELGLR KQTHNNAVVK MLPSFVRRTP DGTENGDFLA LDLGGTNFRV