



P93434Hu01 100μg Myosin IE (MYO1E)

Organism: Homo sapiens (Human)

Instruction manual

kDa

66.2

33

26

20

14.4

15% SDS-PAGE

FOR IN VITRO USE AND RESEARCH USE ONLY
NOT FOR USE IN CLINICAL DIAGNOSTIC PROCEDURES

6th Edition (Revised in March, 2013)

[PROPERTIES]

Residues: Lys781~Pro1036 (Accession # Q12965),

with N-terminal His-Tag.

Host: E. coli

Subcellular Location: Cytoplasm. Cytoplasm,

cytoskeleton. Cytoplasmic vesicle

Clathrin-coated vesicle. Cell junction.

Purity: >95%

Endotoxin Level: <1.0EU per $1\mu g$ (determined by the LAL method).

Formulation: Supplied as lyophilized form in PBS,

pH7.4, containing 5% sucrose, 0.01% sarcosyl.

Predicted isoelectric point: 10.0

Predicted Molecular Mass: 30.1kDa

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

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

[<u>USAGE</u>]

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

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





[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 N-terminal His-Tag, its sequence is listed below. MGHHHHHHSGS-KRDLLLTPKC LYLIGREKVK QGPDKGLVKE VLKRKIEIER ILSVSLSTMQ DDIFILHEQE YDSLLESVFK TEFLSLLAKR YEEKTQKQLP LKFSNTLELK LKKENWGPWS AGGSRQVQFH QGFGDLAVLK PSNKVLQVSI GPGLPKNSRP TRRNTTQNTG YSSGTQNANY PVRAAPPPPG YHQNGVIRNQ YVPYPHAPGS QRSNQKSLYT SMARPPLPRQ QSTSSDRVSQ TPESLDFLKV PDQGAAGVRR QTTSRP