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YBB062Hu02 100μg

Recombinant Heat Shock 70kDa Protein 1 Like Protein (HSPA1L)

Organism Species: Homo sapiens (Human)

Instruction manual

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

9th Edition (Revised in Jul, 2013)

[PROPERTIES]

Residues: Met1~Asp641 (Accession # P34931), with two N-terminal Tags, His-tag and S-tag.

Host: *E. coli*

Purity: >95%

Endotoxin Level: <1.0EU per 1μg

(determined by the LAL method).

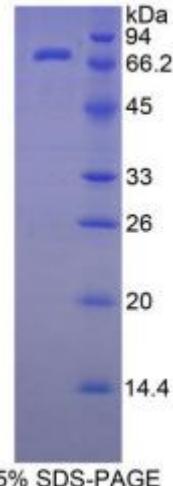
**Formulation: Supplied as lyophilized form in PBS, pH7.4,
containing 5% sucrose, 0.01% sarcosyl.**

Predicted isoelectric point: 5.7

Predicted Molecular Mass: 76.1kDa

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

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



[USAGE]

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



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[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 S-tag, its sequence is listed below.

MHHHHHHSSG LVPRGSGMKE TAAAKFERQH MDSPDLGTDD DDKAMADIGS EF-MATAKGIAIG IDLGTTYSCL GVFQHGKVEI IANDQGNRTT PSYVAFTDTE RLIGDAAKNQ VAMNPQNTVF DAKRLIGRKF NDPVVQADMK LWPFQVINEG GKPVLVSYK GENKAFYPEE ISSMVLTKLK ETAEAFLGHP VTNAVITVPA YFNDSQRQAT KDAGVIAGLN VLRIINEPTA AAIAYGLDKG GQGERHVLIF DLGGGTFDVS ILTIDDGIFE VKATAGDTHL GGEDFDNRDV SHFVEEFKRK HKKDDISQNKR AVRRLRTACE RAKRTLSSST QANLEIDSLY EGIDFYTSIT RARFEELCAD LFRGTLGPVE KALRDAKMDK AKIH DIVLVG GSTRIPKVQR LLQDYFNGRD LNKSINPDEA VAYGAAVQAA ILMGDKSEKV QDLLLLDVAP LSLGLETAGG VMTALIKRNS TIPTKQTQIF TTYSNDNQPGV LIQVYEGERA MTKDNLLGR FDLTGIPPAP RGVPQIEVTF DIDANGILNV TATDKSTGKV NKITITNDKG RLSKEEIERM VLDAEKYKAE DEVQREKIAA KNALESYAFN MKSVVSDEGL KGKISESDKN KILDKCNELL SWLEVNLQAE KDEFDHKRKE LEQMCNPIIT KLYQGGCTGP ACGTGYPGR PATGPTIEEV D