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## YB83541Hu01 100µg Zeta Chain Associated Protein Kinase 70kDa (zAP70)

Organism: Homo sapiens (Human)

Instruction manual

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

7th Edition (Revised in May, 2013)

kDa

94 66.2

45

33

26

20

15% SDS-PAGE

14.4

## [PROPERTIES]

**Residues:** Leu337~Leu600 (Accession # P43403), with two N-terminal Tags, His-tag and T7-tag.

Host: E. coli

Subcellular Location: Cytoplasm. Cell membrane; Peripheral

membrane protein.

**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: 8.6

Predicted Molecular Mass: 34.2kDa

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.

[ 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]

Email:shybio@126.com

The target protein is fused with two N-terminal Tags, His-tag and T7-tag, its sequence is listed below.

MG S S H H H H H H SS G LV P R G S H MA S M T G G Q Q M G R G S E F - L L I A DI E L G C G N F G SVRQGVYRMR KKQIDVAIKV LKQGTEKADT EEMMREAQIM HQLDNPYIVR LIGVCQAEAL MLVMEMAGGG PLHKFLVGKR EEIPVSNVAE LLHQVSMGMK YLEEKNFVHR DLAARNVLLV NRHYAKISDF GLSKALGADD SYYTARSAGK WPLKWYAPEC INFRKFSSRS DVWSYGVTMW EALSYGQKPY KKMKGPEVMA FIEQGKRMEC PPECPPELYA LMSDCWIYKW EDRPDFLTVE QRMRACYYSL