**RPG610Hu01 100µg**

**Recombinant Arylsulfatase B (ARSB)**

**Organism Species: Homo sapiens (Human)**

*Instruction manual*

FOR IN VITRO USE AND RESEARCH USE ONLY

NOT FOR USE IN CLINICAL DIAGNOSTIC PROCEDURES

10th Edition (Revised in Jan, 2014)

# [ PROPERTIES ]
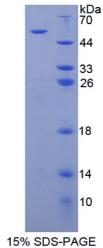
**Residues:** Leu81~Met533

**Tags:** Two N-terminal Tags, His-tag and T7-tag

**Accession:** P15848

**Host:** *E. coli*

**Subcellular Location:** Lysosome.

**Purity:** >90%

**Endotoxin Level:** <1.0EU per 1µg

(determined by the LAL method).

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

**Predicted isoelectric point:** 7.7

**Predicted Molecular Mass:** 55.2kDa

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

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



kDa
70
44
33
26
18
14
10

15% SDS-PAGE

# [ 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 ]

The sequence of the target protein is listed below.

LLDNYYTQPL CTPSRSQLLT GRYQIRTGLQ HQIIWPCQPS CVPLDEKLLP QLLKEAGYTT
HMVGKWHLGM YRKECLPTRR GFDTYFGYLL GSEDYYSHER CTLIDALNVT RCALDFRDGE
EVATGYKNMY STNIFTKRAI ALITNHPPEK PLFLYLALQS VHEPLQVPEE YLKPYDFIQD
KNRHHYAGMV SLMDEAVGNV TAALKSSGLW NNTVFIFSTD NGGQTLAGGN
NWPLRGRKWS LWEGGVRGVG FVASPLLKQK GVKNRELIHI SDWLPTLVKL ARGHTNGTKP
LDGFDVWKTI SEGSPSPRIE LLHNIDPNFV DSSPCPRNSM APAKDDSSLP EYSAFNTSVH
AAIRHGNWKL LTGYPGCGYWFPPPSQYNVS EIPSSDPPTK TLWLFDIDRD PEERHDLSRE
YPHIVTKLLS RLQFYHKHSV PVYFPAQDPR CDPKATGVWG PWM