RPE781Hu01 100ug
Recombinant Annexin A11 (ANXA11)
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: Met1~Asp505
Tags: Two N-terminal Tags, His-tag and T7-tag
Accession: P50995
Host: E. coli
Subcellular Location: Cytoplasm, Melanosome,
Nucleus envelope, Nucleus, nucleoplasm, cytoskeleton, spindle.
Purity: >95\%
Endotoxin Level: <1.0EU per $1 \mu \mathrm{~g}$
(determined by the LAL method).
Formulation: Supplied as lyophilized form in PBS,

pH7.4, containing $5 \%$ trehalose, $0.01 \%$ sarcosyl.
Predicted isoelectric point: 7.5
Predicted Molecular Mass: 58.1 kDa
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, $\mathrm{pH} 7.2-\mathrm{pH} 7.4$.

## [ STORAGE AND STABILITY ]

## Storage: Avoid repeated freeze/thaw cycles.

Store at $2-8^{\circ} \mathrm{C}$ for one month.
Aliquot and store at $-80^{\circ} \mathrm{C}$ 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 $37^{\circ} \mathrm{C}$ for 48 h , 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.
MSYPGYPPPP GGYPPAAPGG GPWGGAAYPP PPSMPPIGLD NVATYAGQFN QDYLSGMAAN MSGTFGGANM PNLYPGAPGA GYPPVPPGGF GQPPSAQQPV PPYGMYPPPG GNPPSRMPSY PPYPGAPVPG QPMPPPGQQP PGAYPGQPPV TYPGQPPVPL PGQQQPVPSSY PGYPGSGTVT PAVPPTQFGS RGTITDAPGF DPLRDAEVLR KAMKGFGTDE QAIIDCLGSR SNKQRQQILL SFKTAYGKDL IKDLKSELSG NFEKTILALM KTPVLFDIYE IKEAIKGVGT DEACLIEILA SRSNEHIREL NRAYKAEFKK TLEEAIRSDT SGHFQRLLIS LSQGNRDEST NVDMSLAQRD AQELYAAGEN RLGTDESKFN AVLCSRSRAH LVAVFNEYQR MTGRDIEKSI CREMSGDLEE GMLAVVKCLK NTPAFFAERL NKAMRGAGTK DRTLIRIMVS RSETDLLDIR SEYKRMYGKS LYHDISGDTS GDYRKILLKI CGGND

