RPB297Hu01 100 $\mu \mathrm{g}$
Recombinant Cytovillin (CVL)
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~Arg295 (Accession \# P15311), with two N-terminal Tags, His-tag and T7-tag.
Host: E. coli
Subcellular Location: Apical cell membrane;
Peripheral membrane protein; Cytoplasmic side.
Cell projection. Microvillus membrane; Ruffle kDa
membrane; Peripheral membrane protein; 18

Cytoplasm, cell cortex. Cytoskeleton.

Purity: >95\%
14
Endotoxin Level: <1.0EU per $1 \mu \mathrm{~g}$
(determined by the LAL method).

Formulation: Supplied as lyophilized form in PBS,
pH7.4, containing $5 \%$ sucrose, $0.01 \%$ sarcosyl.
Predicted isoelectric point: 9.0
Predicted Molecular Mass: 38.6 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, pH7.2-pH7.4.

## [ 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 target protein is fused with two N -terminal Tags, His-tag and T7-tag, its sequence is listed below.
MGSSHHHHHH SSGLVPRGSH MASMTGGQQM GRGSEF- MPKPINVRVT TMDAELEFAI QPNTTGKQLF DQVVKTIGLR EVWYFGLHYV DNKGFPTWLK LDKKVSAQEV RKENPLQFKF RAKFYPEDVA EELIQDITQK LFFLQVKEGI LSDEIYCPPE TAVLLGSYAV QAKFGDYNKE VHKSGYLSSE RLIPQRVMDQ HKLTRDQWED RIQVWHAEHR GMLKDNAMLE YLKIAQDLEM YGINYFEIKN KKGTDLWLGV DALGLNIYEK DDKLTPKIGF PWSEIRNISF NDKKFVIKPI DKKAPDFVFY APRLRINKRI LQLCMGNHEL YMRRR

