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

9th Edition (Revised in Jul, 2013)

[PROPERTIES]

Residues: Val421~Lys629 (Accession # P16284), with N-terminal His-Tag.

Host: *E. coli*

Subcellular Location: Membrane; Single-pass type I membrane protein. Cell junction.

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: 8.8

Predicted Molecular Mass: 24.7kDa

kDa
70
44
33
26
22
18
14
10

15% SDS-PAGE

Accurate Molecular Mass: 18kDa as determined by SDS-PAGE reducing conditions.

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

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

Note: The possible reasons that the actual band size differs from the predicted are as follows:

1. Splice variants: Alternative splicing may create different sized proteins from the same gene.

2. Relative charge: The composition of amino acids may affects the charge of the protein.

3. Post-translational modification: Phosphorylation, glycosylation, methylation etc.

4. Post-translation cleavage: Many proteins are synthesized as pro-proteins, and then cleaved to give the active form.

5. Polymerization of the target protein: Dimerization, multimerization etc.
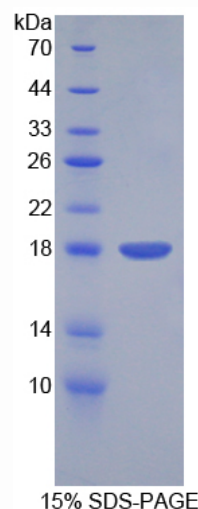
[USAGE]

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 target protein is fused with N-terminal His-Tag, its sequence is listed below.

MGHHHHHHSGSEF- VIKGQTIEVR CESISGTLPI SYQLLKTSKV LENSTKNSND
PAVFKDNPTE DVEYQCVADN CHSHAKMLSE VLRVKVIAPV DEVQISILSS KVVESGEDIV
LQCAVNEGSG PITYKFYREK EGKPFYQMTS NATQAFWTKQ KASKEQEGEY YCTAFNRANH
ASSVPRSKIL TVRVILAPWK KGLIAVVIIG VIIALLIIAA KCYFLRKAK