RPD909Hu01 100 $\mu \mathrm{g}$
Recombinant Dipeptidyl Peptidase 7 (DPP7) Organism Species: Homo sapiens (Human)

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

## [ PROPERTIES ]

Residues: Asp28~Gly486
Tags: Two N-terminal Tags, His-tag and T7-tag
Accession: Q9UHL4
Host: E. coli
Subcellular Location: Lysosome. Cytoplasmic vesicle. Secreted.

Purity: >90\%
kDa70
44332622 pH7.4, containing 5\% trehalose, $0.01 \%$ sarcosyl.
Endotoxin Level: <1.0EU per $1 \mu \mathrm{~g}$ (determined by the LAL method).
Formulation: Supplied as lyophilized form in PBS,

Predicted isoelectric point: 5.5
Predicted Molecular Mass: 54.7 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 sequence of the target protein is listed below.
DPG FQERFFQQRL DHFNFERFGN KTFPQRFLVS DRFWVRGEGP IFFYTGNEGD VWAFANNSAF VAELAAERGA LLVFAEHRYY GKSLPFGAQS TQRGHTELLT VEQALADFAE LLRALRRDLG AQDAPAIAFG GSYGGMLSAY LRMKYPHLVA GALAASAPVL AVAGLGDSNQ FFRDVTADFE GQSPKCTQGV REAFRQIKDL FLQGAYDTVR WEFGTCQPLS DEKDLTQLFM FARNAFTVLA MMDYPYPTDF LGPLPANPVK VGCDRLLSEA QRITGLRALA GLVYNASGSE HCYDIYRLYH SCADPTGCGT GPDARAWDYQ ACTEINLTFA SNNVTDMFPD LPFTDELRQR YCLDTWGVWP RPDWLLTSFW GGDLRAASNI IFSNGNLDPW AGGGIRRNLS ASVIAVTIQG GAHHLDLRAS HPEDPASVVE ARKLEATIIG EWVKAARREQ QPALRG

