#### RPF219Mu01 10µg Recombinant Ectonucleotide Pyrophosphatase/Phosphodiesterase 1 (ENPP1) Organism Species: Mus musculus (Mouse) Instruction manual

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

9th Edition (Revised in Jul, 2013)

### [PROPERTIES]

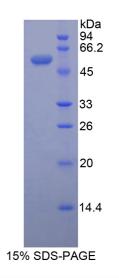
Residues: Gly568~Asp789 (Accession # P06802), with two N-terminal Tags, His-tag and GST-tag. Host: *E. coli* Subcellular Location: Cell membrane; Secreted. Single-pass type II membrane protein. Basolateral cell membrane. Purity: >95% Endotoxin Level: <1.0EU per 1µg (determined by the LAL method). Formulation: Supplied as lyophilized form in 20mM Tris, 500mM NaCl, pH8.0, containing 1mM EDTA, 1mM DTT, 0.01% sarcosyl, 5% trehalose, and preservative. Predicted isoelectric point: 6.4 Predicted Molecular Mass: 58.3kDa Applications: SDS-PAGE; WB; ELISA; IP.

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

# [USAGE]

Reconstitute in ddH<sub>2</sub>O.

Designed by Cloud-Clone Corp., Assembled by Uscn Life Science Inc. ISO9001:2008: ISO13485:20 11271 Richmond Avenue Suite H104, Houston, TX 77982, USA | Toil free: 001-888-960-7402 | Fax: 001-832-538-0088 | Http://www.cloud-clone.us | E-mail: mail@cloud-clo Export Processing Zone Building F, Wuhan, Hubel 430056, PRC | Toil free: 0086-800-880-0687 | Fax: 0086-27-8425-9551 | Http://www.uscnk.com | E-mail: mail@cloud-clo





# [ 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.

# [<u>SEQUENCES</u>]

The target protein is fused with two N-terminal Tags, His-tag and GST-tag, its sequence is listed below.

MSPILGYWKI KGLVQPTRLL LEYLEEKYEE HLYERDEGDK WRNKKFELGL EFPNLPYYID GDVKLTQSMA IIRYIADKHN MLGGCPKERA EISMLEGAVL DIRYGVSRIA YSKDFETLKV DFLSKLPEML KMFEDRLCHK TYLNGDHVTH PDFMLYDALD VVLYMDPMCL DAFPKLVCFK KRIEAIPQID KYLKSSKYIA WPLQGWQATF GGGDHPPKSD GSTSGSGHHH HHHSAGLVPR GSTAIGMKET AAAKFERQHM DSPDLGTGGG SGIEGRGSMG YRGSEF-GSH GSLNHLLKKP IYNPSHPKEE GFLSQCPIKS TSNDLGCTCD PWIVPIKDFE KQLNLTTEDV DDIYHMTVPY GRPRILLKQH HVCLLQQQQF LTGYSLDLLM PLWASYTFLR NDQFSRDDFS NCLYQDLRIP LSPVHKCSYY KSNSKLSYGF LTPPRLNRVS NHIYSEALLT SNIVPMYQSF QVIWHYLHDT LLQRYAHERN GINVVSGPVF DFDYDGRYD

# [REFERENCES]

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