

RPG932Ra01 100ug

Recombinant UDP Glucuronosyltransferase 2 Family, Polypeptide B7 (UGT2B7)

Organism Species: Rattus norvegicus (Rat)

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: Phe18~Val495

Tags: Two N-terminal Tags, His-tag and T7-tag

Accession: Q62789

Host: *E. coli*

Subcellular Location: Microsome membrane;
Single-pass membrane protein. Endoplasmic
reticulum membrane.

Purity: >95%

Endotoxin Level: <1.0EU per 1µg
(determined by the LAL method).

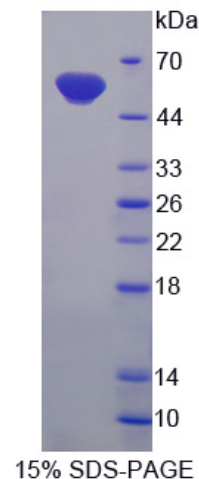
Formulation: Supplied as lyophilized form in 20mM Tris,
150mM NaCl, pH8.0, containing 1mM EDTA, 1mM DTT,
0.01% sarcosyl, 5% trehalose, and preservative.

Predicted isoelectric point: 7.0

Predicted Molecular Mass: 57.8kDa

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 ddH₂O.

[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 sequence of the target protein is listed below.

FRS GNCGKVLVWP LEYSHWMNLK IILDELVQRG HEVTVLRPSS SVSLDPKKAS
GLVYETSPTT SNNDEVEKSF YPVGDMWTYD VPKYTCLRYY PSLNKMFGQF SDLWLQLCRE
VVSNKELIAK LKESQFDVVL SDAVGPCGEL IAEILQLPFV YSLRFATAPG IEKYSAGQPF
PPSYVPIILS GFSGQMTFME RVENMLCLLY FDSWFESFPA KDWDPFSEI LGRPTTMVDT
MKKAEIWLIR SYWDLEFPRP SLPNIEFVGG LHCQPAKPLP KEMEDFAQSS GEHGVVWFSL
GSMIRNITQE RANTIASALA QIPQKVFWRP EGKKPDTLGP NTRVFKWIPQ NDLLGHPKTK
AFVTHGGANG IYESIHYGIP PMVGIPLFAE QRDNVAHMVA KGAAVSIDFH TMSSDLLNA
LKAVINNPST KKKVMWLSAI HHDQPLKPLD RAVFWIEFVM RHKGAKHLRP LAHNLALVSV
HSLDV