

RPB766Mu01 100µg

Recombinant Lectin Galactoside Binding, Soluble 3 Binding Protein (LGALS3BP)

Organism Species: *Mus musculus* (Mouse)

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: Thr19~Val577

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

Accession: Q07797

Host: *E. coli*

Subcellular Location: Secreted, extracellular space, extracellular matrix.

Purity: >90%

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

Formulation: Supplied as lyophilized form in PBS, pH7.4, containing 5% trehalose, 0.01% sarcosyl.

Predicted isoelectric point: 5.0

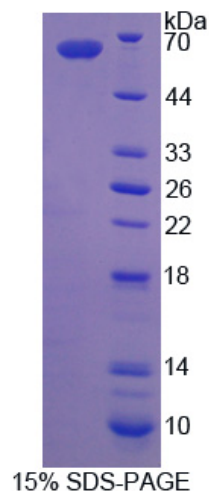
Predicted Molecular Mass: 66.0kDa

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°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.

TE DGDMRLVNGA SANEGRVEIF YRGRWGTVCD NLWNLLDAHV VCRALGYENA
TQALGRAAFG PGKGPIMLDE VECTGTESSL ASCRSLGWMV SRCGHEKDAG
VVCSDNTTGL HILDLSGELS DALGQIFDSQ QGCDLFIQVT GQGYESLCLC AHTLILRTNP
EAQALWQVVG SSVIMRVDAE CMPVVRDFLR YFYRRRIEVS MSSVKCLHKL ASAYGATELQ
DYCGRLFATL LPQDPTFHTP LDLYAYARAT GDSMLEDLCV QFLAWNFEPL TQSESWSAVP
TTLIQALLPK SELAVSSELD LLKAVDQWST ETIASHEDIE RLVEQVRFPMLPQELFELQ
FNLSLYQDHFQ ALFQRKTMQA LEFHTVPVEV LAKYKGLNLT EDYKPRPLYT SSTWSSLVMA
STWRAQRYEY NRYNQLYTYG YGSVARYNSY QSFQTPQHPS FLFKDKQISW SATYLPTMQS
CWNYGFSCTS NELPVLGLTT SSYSNPTIGY ENRVLILCGG YSVVDVTSFE GSKAPIPTAL
DTNSSKTPSL FPCASGAFSS FRVVIRPFYL TNSTDMV