RPH136Mu01 100ug

Recombinant Serine Palmitoyltransferase, Long Chain Base Subunit 3 (SPTLC3)

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: Arg80~Ala563

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

Accession: Q8BG54

Host: *E. coli*

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

Purity: >90%

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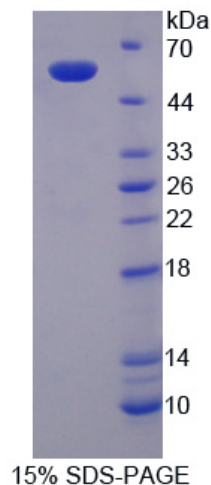
(determined by the LAL method).

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: 7.3

Predicted Molecular Mass: 58.4kDa

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

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

kDa
70
44
33
26
22
18
14
10

15% SDS-PAGE

[USAGE]

Reconstitute in sterile ddH$_2$O.

[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.

R DFMRNWGIEK CNAAVEREEQ KDFVPLYQDF ENFYKRNLYM RIRDSWSHTV
CSAPEPYMNV MEKVTDDYNW TFRHTGKVIE NIINMASYNY LGLAGKYDDS MVRVKDTLEK
YGVGVASTRN EMGTLDIHKE LEDLMAEFLN VEAVMSFGMG FATNAMNIPV FVGKGCLILS
DEFNHTSVIL GSRLSGAVIR PFKHNNAENL EKLLREAIIR GQPGTGRAWK KILIVVEGVY
SMEGSIVNLA QIVALKKKYK AYLYIDEAHS IGCTGPTGRG VRELFGLDPE DIDVYMGTFT
KSFSGSGGYI GGKKEIVDYL RMQSHSTTYA TSMSPVVAAQ LIRSLKITMG YEGNIGGMER
IQQLKENIKY FRRRLKEMGF IIYGNDFSPV IPVLLYMPAK VSAFSRFLLK KKISVVVVGF
PATSLPEGRA RFSMSSAHTR EMLDTVLEVV DELGDLLNVK YFPLKKSGRA ILYNKEGFDN
EASFEEMHSE PEA