YBA383Mu01 100µg

Instruction manual

Recombinant Phosphoinositide-3-Kinase Class-2-Alpha Polypeptide (PIK3C2a)

Organism Species: Mus musculus (Mouse)

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

9th Edition (Revised in Jul, 2013)

[PROPERTIES]

Residues: Lys349~Pro611 (Accession # Q61194), with

two N-terminal Tags, His-tag and GST-tag.

Host: E. coli

Subcellular Location: Cell membrane. Golgi

apparatus. Cytoplasmic vesicle, clathrin-coated

vesicle. Nucleus. Cytoplasm.

Purity: >95%

Endotoxin Level: <1.0EU per 1µg

(determined by the LAL method).

Formulation: Supplied as lyophilized form in PBS, pH7.4, 15% SDS-PAGE

containing 5% trehalose, 0.01% sarcosyl.

Predicted isoelectric point: 5.8

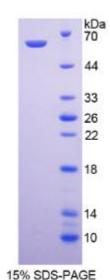
Predicted Molecular Mass: 61.9kDa

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

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

[<u>USAGE</u>]

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 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 GDVKLTOSMA IIRYIADKHN MLGGCPKERA EISMLEGAVL DIRYGVSRIA YSKDFETLKV DFLSKLPEML KMFEDRLCHK TYLNGDHVTH PDFMLYDALD VVLYMDPMCL DAFPKLVCFK KRIEAIPQID KYLKSSKYIA WPLQGWQATF GGGDHPPKSD GSTSGSGHHH HHHSAGLVPR GSTAIGMKET AAAKFERQHM DSPDLGTGGG SGIEGRGSMG YRGSEF-KA QGQVSQKDPN GTSSLPTGSS LLQEFEVQND EVAAFCQSIM KLKTKFPYTD HCTNPGYLLS PVTVQRNMCG ENASVKVSIE IEGLQLPVTF TCDVSSTVEI IIMQALCWVH DDLNQVDVGS YILKVCGQEE VLQNNHCLGS HEHIQNCRKW DTEIKLQLLT LSAMCQNLAR TAEDDEAPVD LNKYLYQIEK PYKEVMTRHP VEELLDSYHY QVELALQTEN QHRAVDQVIK AVRKICSALD GVETPSVTEA VKKLKRAVNL P