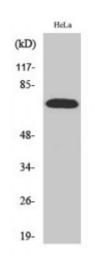


Anti-p70 S6 kinase alpha antibody



Description

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Rabbit polyclonal to p70 S6 kinase alpha.

Model STJ94920

Host Rabbit

Reactivity Human, Mouse, Rat

Applications ELISA, IF, IHC, WB

Immunogen Synthesized peptide derived from human p70 S6 kinase alpha around the non-

phosphorylation site of S371.

Immunogen Region 310-390 aa

Gene ID <u>6198</u>

Gene Symbol RPS6KB1

Dilution range WB 1:500-1:2000IHC 1:100-1:300IF 1:200-1:1000ELISA 1:20000

Specificity p70 S6 kinase alpha Polyclonal Antibody detects endogenous levels of p70 S6

kinase alpha protein.

Tissue Specificity Widely expressed.

Purification The antibody was affinity-purified from rabbit antiserum by affinity-

chromatography using epitope-specific immunogen.

Note For Research Use Only (RUO).

Protein Name Ribosomal protein S6 kinase beta-1 S6K-beta-1 S6K1 70 kDa ribosomal

protein S6 kinase 1 P70S6K1 p70-S6K 1 Ribosomal protein S6 kinase I Serine/threonine-protein kinase 14A p70 ribosomal S6 kinase alpha

Molecular Weight 60 kDa

Clonality Polyclonal

Conjugation Unconjugated

Isotype IgG

Formulation Liquid in PBS containing 50% glycerol, 0.5% BSA and 0.02% sodium azide.

Concentration 1 mg/ml

Storage Instruction Store at -20°C, and avoid repeat freeze-thaw cycles.

Database Links HGNC:10436OMIM:608938

Alternative Names Ribosomal protein S6 kinase beta-1 S6K-beta-1 S6K1 70 kDa ribosomal

protein S6 kinase 1 P70S6K1 p70-S6K 1 Ribosomal protein S6 kinase I Serine/threonine-protein kinase 14A p70 ribosomal S6 kinase alpha

Function Serine/threonine-protein kinase that acts downstream of mTOR signaling in

response to growth factors and nutrients to promote cell proliferation, cell growth and cell cycle progression. Regulates protein synthesis through phosphorylation of EIF4B, RPS6 and EEF2K, and contributes to cell survival by repressing the pro-apoptotic function of BAD. Under conditions of nutrient depletion, the inactive form associates with the EIF3 translation initiation complex. Upon mitogenic stimulation, phosphorylation by the mammalian target of rapamycin complex 1 (mTORC1) leads to dissociation from the EIF3 complex and activation. The active form then phosphorylates and activates several substrates in the pre-initiation complex, including the EIF2B complex and the cap-binding complex component EIF4B. Also controls translation initiation by phosphorylating a negative regulator of EIF4A, PDCD4, targeting it for ubiquitination and subsequent proteolysis. Promotes initiation

of the pioneer round of protein synthesis by phosphorylating

POLDIP3/SKAR. In response to IGF1, activates translation elongation by phosphorylating EEF2 kinase (EEF2K), which leads to its inhibition and thus activation of EEF2. Also plays a role in feedback regulation of mTORC2 by

mTORC1 by phosphorylating RICTOR, resulting in the inhibition of mTORC2 and AKT1 signaling. Mediates cell survival by phosphorylating the pro-apoptotic protein BAD and suppressing its pro-apoptotic function. Phosphorylates mitochondrial URI1 leading to dissociation of a URI1-

PPP1CC complex. The free mitochondrial PPP1CC can then dephosphorylate

RPS6KB1 at Thr-412, which is proposed to be a negative feedback mechanism for the RPS6KB1 anti-apoptotic function. Mediates TNF-alphainduced insulin resistance by phosphorylating IRS1 at multiple serine residues, resulting in accelerated degradation of IRS1. In cells lacking functional TSC1-2 complex, constitutively phosphorylates and inhibits GSK3B. May be involved in cytoskeletal rearrangement through binding to

neurabin. Phosphorylates and activates the pyrimidine biosynthesis enzyme

CAD, downstream of MTOR.

Sequence and Domain Family The autoinhibitory domain is believed to block phosphorylation within the

AGC-kinase C-terminal domain and the activation loop.; The TOS (TOR

signaling) motif is essential for activation by mTORC1.

Cellular Localization Cell junction, synapse, synaptosome Mitochondrion outer membrane.

Mitochondrion. Colocalizes with URI1 at mitochondrion.. Isoform Alpha I:

Nucleus. Cytoplasm.. Isoform Alpha II: Cytoplasm.

Post-translational Modifications

Phosphorylation at Thr-412 is regulated by mTORC1. The phosphorylation at this site is maintained by an agonist-dependent autophosphorylation mechanism . Activated by phosphorylation at Thr-252 by PDPK1. Dephosphorylation by PPP1CC at Thr-412 in mitochondrion.

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