



# **RPS6KA5 Blocking Peptide (C-term)**

Synthetic peptide Catalog # BP21184b

# **Specification**

RPS6KA5 Blocking Peptide (C-term) - Product Information

Primary Accession <u>075582</u>

RPS6KA5 Blocking Peptide (C-term) - Additional Information

**Gene ID 9252** 

#### **Other Names**

Ribosomal protein S6 kinase alpha-5, S6K-alpha-5, 90 kDa ribosomal protein S6 kinase 5, Nuclear mitogen- and stress-activated protein kinase 1, RSK-like protein kinase, RSKL, RPS6KA5, MSK1

# **Target/Specificity**

The synthetic peptide sequence is selected from aa 731-745 of HUMAN RPS6KA5

### **Format**

Peptides are lyophilized in a solid powder format. Peptides can be reconstituted in solution using the appropriate buffer as needed.

#### Storage

Maintain refrigerated at 2-8°C for up to 6 months. For long term storage store at -20°C.

#### **Precautions**

This product is for research use only. Not for use in diagnostic or therapeutic procedures.

RPS6KA5 Blocking Peptide (C-term) - Protein Information

Name RPS6KA5

Synonyms MSK1

## **Function**

Serine/threonine-protein kinase that is

# RPS6KA5 Blocking Peptide (C-term) - Background

Serine/threonine-protein kinase that is required for the mitogen or stress-induced phosphorylation of the transcription factors CREB1 and ATF1 and for the regulation of the transcription factors RELA, STAT3 and ETV1/ER81, and that contributes to gene activation by histone phosphorylation and functions in the regulation of inflammatory genes. Phosphorylates CREB1 and ATF1 in response to mitogenic or stress stimuli such as UV-C irradiation, epidermal growth factor (EGF) and anisomycin. Plays an essential role in the control of RELA transcriptional activity in response to TNF and upon glucocorticoid, associates in the cytoplasm with the glucocorticoid receptor NR3C1 and contributes to RELA inhibition and repression of inflammatory gene expression. In skeletal myoblasts is required for phosphorylation of RELA at 'Ser-276' during oxidative stress. In erythropoietin-stimulated cells, is necessary for the 'Ser-727' phosphorylation of STAT3 and regulation of its transcriptional potential. Phosphorylates ETV1/ER81 at 'Ser-191' and 'Ser-216', and thereby regulates its ability to stimulate transcription, which may be important during development and breast tumor formation. Directly represses transcription via phosphorylation of 'Ser-1' of histone H2A. Phosphorylates 'Ser-10' of histone H3 in response to mitogenics, stress stimuli and EGF, which results in the transcriptional activation of several immediate early genes, including proto- oncogenes c-fos/FOS and c-jun/JUN. May also phosphorylate 'Ser-28' of histone H3. Mediates the mitogen- and stress-induced phosphorylation of high mobility group protein 1 (HMGN1/HMG14). In lipopolysaccharide-stimulated primary macrophages, acts downstream of the Toll-like receptor TLR4 to limit the production of proinflammatory cytokines. Functions probably by inducing transcription of the MAP kinase phosphatase DUSP1 and the antiinflammatory cytokine interleukin 10 (IL10), via





required for the mitogen or stress-induced phosphorylation of the transcription factors CREB1 and ATF1 and for the regulation of the transcription factors RELA, STAT3 and ETV1/ER81, and that contributes to gene activation by histone phosphorylation and functions in the regulation of inflammatory genes (PubMed:<a href="http://www.uniprot.org/citations/11909979""

target="\_blank">11909979</a>,

PubMed:<a href="http://www.uniprot.org/ci tations/12569367"

target="\_blank">12569367</a>,

PubMed: <a href="http://www.uniprot.org/ci tations/12763138"

target=" blank">12763138</a>,

PubMed: <a href="http://www.uniprot.org/ci tations/9687510"

target=" blank">9687510</a>,

PubMed:<a href="http://www.uniprot.org/ci tations/18511904"

target=" blank">18511904</a>,

PubMed:<a href="http://www.uniprot.org/ci tations/9873047"

target=" blank">9873047</a>).

Phosphorylates CREB1 and ATF1 in

response to mitogenic or stress stimuli such as UV-C irradiation, epidermal growth factor (EGF) and anisomycin (PubMed:<a href="http://www.uniprot.org/citations/11909979" target="blank">11909979</a>,

PubMed:<a href="http://www.uniprot.org/ci tations/9873047"

target="\_blank">9873047</a>). Plays an essential role in the control of RELA transcriptional activity in response to TNF and upon glucocorticoid, associates in the cytoplasm with the glucocorticoid receptor NR3C1 and contributes to RELA inhibition and repression of inflammatory gene expression (PubMed:<a href="http://www.u niprot.org/citations/12628924" target=" blank">12628924</a>,

PubMed: <a href="http://www.uniprot.org/ci tations/18511904"

target="\_blank">18511904</a>). In skeletal myoblasts is required for phosphorylation of RELA at 'Ser-276' during oxidative stress (PubMed:<a href="http://www.uniprot.org/citations/12628924" target="\_blank">12628924</a>). In erythropoietin-stimulated cells, is necessary for the 'Ser-727' phosphorylation of STAT3 and regulation of its transcriptional potential (PubMed:<a href="http://www.uniprot.org/citations/12763138" target=" blank">12763138</a>).

CREB1 and ATF1 transcription factors. Plays a role in neuronal cell death by mediating the downstream effects of excitotoxic injury.

# RPS6KA5 Blocking Peptide (C-term) - References

Deak M.,et al.EMBO J. 17:4426-4441(1998). New L.,et al.J. Biol. Chem. 274:1026-1032(1999). Jiang C.,et al.Cytogenet. Cell Genet. 87:261-262(1999). Heilig R.,et al.Nature 421:601-607(2003). Wiggin G.R.,et al.Mol. Cell. Biol. 22:2871-2881(2002).



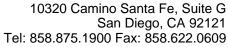
Phosphorylates ETV1/ER81 at 'Ser-191' and 'Ser-216', and thereby regulates its ability to stimulate transcription, which may be important during development and breast tumor formation (PubMed:<a href="http://w ww.uniprot.org/citations/12569367" target=" blank">12569367</a>). Directly represses transcription via phosphorylation of 'Ser-1' of histone H2A (PubMed: <a href=" http://www.uniprot.org/citations/15010469" target=" blank">15010469</a>). Phosphorylates 'Ser-10' of histone H3 in response to mitogenics, stress stimuli and EGF, which results in the transcriptional activation of several immediate early genes, including proto-oncogenes c-fos/FOS and c-jun/JUN (PubMed:<a href="http://ww w.uniprot.org/citations/12773393" target=" blank">12773393</a>). May also phosphorylate 'Ser-28' of histone H3 (PubMed:<a href="http://www.uniprot.org/c itations/12773393" target=" blank">12773393</a>). Mediates the mitogen- and stress-induced phosphorylation of high mobility group protein 1 (HMGN1/HMG14) (PubMed:<a href ="http://www.uniprot.org/citations/1277339 3" target=" blank">12773393</a>). In lipopolysaccharide-stimulated primary macrophages, acts downstream of the Toll-like receptor TLR4 to limit the production of pro-inflammatory cytokines (By similarity). Functions probably by inducing transcription of the MAP kinase phosphatase DUSP1 and the anti-inflammatory cytokine interleukin 10 (IL10), via CREB1 and ATF1 transcription factors (By similarity). Plays a role in neuronal cell death by mediating the downstream effects of excitotoxic injury (By similarity). Phosphorylates TRIM7 at 'Ser-107' in response to growth factor signaling via the MEK/ERK pathway, thereby stimulating its ubiquitin ligase activity (PubMed:<a href="http://www.uniprot.org/c itations/25851810" target=" blank">25851810</a>).

#### **Cellular Location**

Nucleus. Cytoplasm. Note=Predominantly nuclear. Exported into cytoplasm in response to glucocorticoid

#### **Tissue Location**

Widely expressed with high levels in heart, brain and placenta. Less abundant in lung, kidney and liver





# RPS6KA5 Blocking Peptide (C-term) -**Protocols**

Provided below are standard protocols that you may find useful for product applications.

• Blocking Peptides