:6269 Store at -20°C

SignalSilence® p38lpha MAPK siRNA l

10 μM in 300 μl
 (100 transfections)

rev. 02/09/16



Species Cross-Reactivity: H

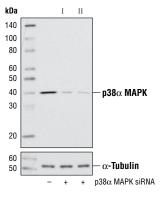
Description: SignalSilence[®] p38 α MAPK siRNA I from Cell Signaling Technology (CST) allows the researcher to specifically inhibit p38 α MAPK expression using RNA interference, a method whereby gene expression can be selectively silenced through the delivery of double stranded RNA molecules into the cell. All SignalSilence[®] siRNA products from CST are rigorously tested in-house and have been shown to reduce target protein expression by western analysis.

Background: p38 MAP kinase (MAPK), also called RK (1) or CSBP (2), is the mammalian orthologue of the yeast HOG kinase that participates in a signaling cascade controlling cellular responses to cytokines and stress (1-4). Four isoforms of p38 MAP kinase, p38α, β, γ (also known as ERK6 or SAPK3) and δ (also known as SAPK4) have been identified. Similar to the SAPK/JNK pathway, p38 MAP kinase is activated by a variety of cellular stresses including osmotic shock, inflammatory cytokines, lipopolysaccharides (LPS), UV light and growth factors (1-5). MKK3, MKK6 and SEK activate p38 MAP kinase by phosphorylation at Thr180 and Tyr182. Activated p38 MAP kinase has been shown to phosphorylate the transcription factors ATF-2 (5), Max (6) and MEF2 (5-8).

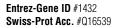
SB203580 (4-(4-fluorophenyl)-2-(4-methylsulfinylphenyl)-5-(4- pyridyl)imidazole) is a selective inhibitor of p38 MAPK. This compound inhibits the activation of MAPKAPK-2 by p38 and thereby phosphorylation of HSP27 (9). SB203580 inhibits p38 catalytic activity by binding to the ATP binding pocket, but does not inhibit phosphorylation of p38 by upstream kinases (10).

Directions for Use: CST recommends transfection with 100 nM p38 α MAPK siRNA I 48 to 72 hours prior to cell lysis. For transfection procedure, follow protocol provided by the transfection reagent manufacturer. Please feel free to contact CST with any questions on use.

Quality Control: Oligonucleotide synthesis is monitored base by base through trityl analysis to ensure appropriate coupling efficiency. The oligo is subsequently purified by affinity-solid phase extraction. The annealed RNA duplex is further analyzed by mass spectrometry to verify the exact composition of the duplex. Each lot is compared to the previous lot by mass spectrometry to ensure maximum lot-to-lot consistency.



Western blot analysis of extracts from 293 cells, transfected with 100 nM SignalSilence® Control siRNA (Unconjugated) #6568 (-), SignalSilence® p38 α MAPK siRNA I (+) or SignalSilence® p38 α MAPK siRNA II #6277 (+), using p38 α MAPK Antibody #9218 (upper) or α -Tubulin (11H10) Rabbit mAb #2125 (lower). The p38 α MAPK Antibody confirms silencing of p38 α MAPK expression, while the α -Tubulin (11H10) Rabbit mAb is used as a loading control.



Storage: $p38\alpha$ MAPK siRNA I is supplied in RNAse-free water. *Aliquot and store at -20°C.*

Please visit www.cellsignal.com for a complete listing of recommended companion products.

Background References:

(1) Rouse, J. et al. (1994) Cell 78, 1027-1037.

Cell Signaling

Orders 877-616-CELL (2355)

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- (2) Han, J. et al. (1994) Science 265, 808-811.
- (3) Lee, J.C. et al. (1994) Nature 372, 739-746.
- (4) Freshney, N.W. et al. (1994) Cell 78, 1039-1049.
- (5) Raingeaud, J. et al. (1995) J. Biol. Chem. 270, 7420-7426.
- (6) Zervos, A.S. et al. (1995) Proc. Natl. Acad. Sci. USA 92, 10531-10534.
- (7) Zhao, M. et al. (1999) Mol. Cell. Biol. 19, 21-30.
- (8) Yang, S.H. et al. (1999) Mol. Cell. Biol. 19, 4028-4038.

 Applications Key:
 W—Western
 IP—Immunoprecipitation
 IHC—Immunohistochemistry
 ChIP—Chromatin Immunoprecipitation
 IF—Immunofluorescence
 F—Flow cytometry
 E-P-ELISA-Peptide

 Species Cross-Reactivity Key:
 H—human
 M—mouse
 R—rat
 Hm—hamster
 Mk—monkey
 Mi—mink
 C—chicken
 Dm—D. melanogaster
 X—zebrafish
 B—bovine

 Dg—dog
 Pg—pig
 Sc—S. cerevisiae
 Ce-C. elegans
 Hr—Horse
 AII—all species expected
 Species enclosed in parentheses are predicted to react based on 100% homology.