SANTA CRUZ BIOTECHNOLOGY, INC.

c-Myb siRNA (h): sc-29855



BACKGROUND

The highly leukemogenic avian retrovirus E26 contains two oncogenes, v-Myb and v-Ets, which are expressed together as a fusion protein. The cellular homolog of v-Myb, designated c-Myb, encodes a transcription factor. Deletion or disruption of a negative regulatory domain mapping within the carboxy terminal domain of c-Myb results in enhanced transactivating capacity and in parallel, leads to activation of its ability to transform hemopoietic cells. c-Myb is expressed preferentially, but not exclusively, in immature hemopoietic cells and its expression decreases as cells differentiate. A second member of the Myb proto-oncogene family, B-Myb, encodes a second sequence-specific DNA binding protein. B-Myb RNA levels are low or undetectable in quiescent cells but increase at the G_1 to S phase transition following mitogenic stimulation. Studies suggest that B-Myb expression rescues cells from p53-induced G_1 arrest mediated by p21.

REFERENCES

- Gonda, T.J., et al. 1984. Expression of Myb, Myc and Fos proto-oncogenes during the differentiation of a murine myeloid leukaemia. Nature 310: 249-251.
- Gonda, T.J., et al. 1985. Nucleotide sequence of cDNA clones of the murine Myb proto-oncogene. EMBO J. 4: 2003-2008.
- Sakura, H., et al. 1989. Delineation of three functional domains of the transcriptional activator encoded by the c-Myb protooncogene. Proc. Natl. Acad. Sci. USA 86: 5758-5762.
- Mizuguchi, G., et al. 1990. DNA binding activity and transcriptional activator function of the human B-Myb protein compared with c-Myb. J. Biol. Chem. 265: 9280-9284.

CHROMOSOMAL LOCATION

Genetic locus: MYB (human) mapping to 6q23.3.

PRODUCT

c-Myb siRNA (h) is a pool of 3 target-specific 19-25 nt siRNAs designed to knock down gene expression. Each vial contains 3.3 nmol of lyophilized siRNA, sufficient for a 10 μ M solution once resuspended using protocol below. Suitable for 50-100 transfections. Also see c-Myb shRNA Plasmid (h): sc-29855-SH and c-Myb shRNA (h) Lentiviral Particles: sc-29855-V as alternate gene silencing products.

For independent verification of c-Myb (h) gene silencing results, we also provide the individual siRNA duplex components. Each is available as 3.3 nmol of lyophilized siRNA. These include: sc-29855A, sc-29855B and sc-29855C.

STORAGE AND RESUSPENSION

Store lyophilized siRNA duplex at -20° C with desiccant. Stable for at least one year from the date of shipment. Once resuspended, store at -20° C, avoid contact with RNAses and repeated freeze thaw cycles.

Resuspend lyophilized siRNA duplex in 330 μ l of the RNAse-free water provided. Resuspension of the siRNA duplex in 330 μ l of RNAse-free water makes a 10 μ M solution in a 10 μ M Tris-HCl, pH 8.0, 20 mM NaCl, 1 mM EDTA buffered solution.

APPLICATIONS

c-Myb siRNA (h) is recommended for the inhibition of c-Myb expression in human cells.

SUPPORT REAGENTS

For optimal siRNA transfection efficiency, Santa Cruz Biotechnology's siRNA Transfection Reagent: sc-29528 (0.3 ml), siRNA Transfection Medium: sc-36868 (20 ml) and siRNA Dilution Buffer: sc-29527 (1.5 ml) are recommended. Control siRNAs or Fluorescein Conjugated Control siRNAs are available as 10 μ M in 66 μ l. Each contain a scrambled sequence that will not lead to the specific degradation of any known cellular mRNA. Fluorescein Conjugated Control siRNAs include: sc-36869, sc-44239, sc-44240 and sc-44241. Control siRNAs include: sc-37007, sc-44230, sc-44231, sc-44232, sc-44233, sc-44234, sc-44235, sc-44236, sc-44237 and sc-44238.

GENE EXPRESSION MONITORING

c-Myb (D-7): sc-74512 is recommended as a control antibody for monitoring of c-Myb gene expression knockdown by Western Blotting (starting dilution 1:200, dilution range 1:100-1:1000) or immunofluorescence (starting dilution 1:50, dilution range 1:50-1:500).

RT-PCR REAGENTS

Semi-quantitative RT-PCR may be performed to monitor c-Myb gene expression knockdown using RT-PCR Primer: c-Myb (h)-PR: sc-29855-PR (20 μ l, 590 bp). Annealing temperature for the primers should be 55-60° C and the extension temperature should be 68-72° C.

SELECT PRODUCT CITATIONS

- Tapias, A., et al. 2008. Transcriptional regulation of the 5'-flanking region of the human transcription factor Sp3 gene by NF-1, c-Myb, B-Myb, AP-1 and E2F. Biochim. Biophys. Acta 1779: 318-329.
- Yang, H., et al. 2011. Insulin-like growth factor 1 activates methionine adenosyltransferase 2A transcription by multiple pathways in human colon cancer cells. Biochem. J. 436: 507-516.
- Vargova, K., et al. 2011. MYB transcriptionally regulates the miR-155 host gene in chronic lymphocytic leukemia. Blood 117: 3816-3825.
- Basova, P., et al. 2014. Aggressive acute myeloid leukemia in PU.1/p53 double-mutant mice. Oncogene 33: 4735-4745.
- Kodric, K., et al. 2019. Sex-determining region Y (SRY) attributes to gender differences in RANKL expression and incidence of osteoporosis. Exp. Mol. Med. 51: 97.
- Zhang, X.Y., et al. 2020. Regulation of Myb mediated cisplatin resistance of ovarian cancer cells involves miR-21-Wnt signaling axis. Sci. Rep. 10: 6893.
- Kaundal, B., et al. 2020. Nanoformulation of EPZ011989 attenuates EZH2c-Myb epigenetic interaction by proteasomal degradation in acute myeloid leukemia. Mol. Pharm. 17: 604-621.

RESEARCH USE

For research use only, not for use in diagnostic procedures.