

CHD1L siRNA (m): sc-105201

BACKGROUND

CHD1L (chromodomain helicase DNA-binding protein 1-like) is an 897 amino acid protein encoded by the human gene CHD1L. The CHD family of proteins are ATP-dependent chromatin remodeling enzymes which combine chromodomains with SWI2/SNF2 ATPase/helicase motifs and DNA-binding capability. Chromodomains are protein regions of approximately 40-50 amino acid residues that are found in proteins associated with chromatin remodeling and manipulation. The domain is highly conserved among both plants and animals and is found in a large variety of proteins from many genomes. CHD1L and CHD1 share most homology within two internal helicase domains.

REFERENCES

1. Stokes, D.G. and Perry, R.P. 1995. DNA-binding and chromatin localization properties of CHD1. *Mol. Cell. Biol.* 15: 2745-2753.
2. Woodage, T., Basrai, M.A. and Baxevanis, A.D. 1997. Characterization of the CHD family of proteins. *Proc. Natl. Acad. Sci. USA* 94: 11472-11477.
3. Kelley, D.E., Stokes, D.G. and Perry, R.P. 1999. CHD1 interacts with SSRP1 and depends on both its chromodomain and its ATPase/helicase-like domain for proper association with chromatin. *Chromosoma* 108: 10-25.
4. Tai, H.H., Geisterfer, M. and Bell, J.C. 2003. CHD1 associates with NCoR and histone deacetylase as well as with RNA splicing proteins. *Biochem. Biophys. Res. Commun.* 308: 170-176.
5. Sims, R.J., Chen, C.F. and Santos-Rosa, H. 2006. Human but not yeast CHD1 binds directly and selectively to Histone H3 methylated at Lysine 4 via its tandem chromodomains. *J. Biol. Chem.* 280: 41789-41792.
6. Flanagan, J.F., Mi, L.Z. and Chruszcz, M. 2006. Double chromodomains cooperate to recognize the methylated Histone H3 tail. *Nature* 438: 1181-1185.

CHROMOSOMAL LOCATION

Genetic locus: Chd1l (mouse) mapping to 3 F2.2.

PRODUCT

CHD1L siRNA (m) 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 CHD1L shRNA Plasmid (m): sc-105201-SH and CHD1L shRNA (m) Lentiviral Particles: sc-105201-V as alternate gene silencing products.

For independent verification of CHD1L (m) gene silencing results, we also provide the individual siRNA duplex components. Each is available as 3.3 nmol of lyophilized siRNA. These include: sc-105201A, sc-105201B and sc-105201C.

RESEARCH USE

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

PROTOCOLS

See our web site at www.scbt.com for detailed protocols and support products.

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

CHD1L siRNA (m) is recommended for the inhibition of CHD1L expression in mouse 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.

RT-PCR REAGENTS

Semi-quantitative RT-PCR may be performed to monitor CHD1L gene expression knockdown using RT-PCR Primer: CHD1L (m)-PR: sc-105201-PR (20 μ l). Annealing temperature for the primers should be 55-60° C and the extension temperature should be 68-72° C.