

HoxC5 siRNA (h): sc-75287

BACKGROUND

The Hox homeobox genes encode proteins that are transcriptional regulators with an established role in embryonic development. HoxC5 (homeobox C5), also known as CP11, HOX3 or HOX3D, is a 222 amino acid protein that localizes to the nucleus and contains one homeobox DNA-binding domain. Existing as multiple alternatively spliced isoforms, HoxC5 functions as a sequence-specific DNA-binding transcription factor that is part of a regulatory mechanism that provides cells with positional identities during development. The gene encoding HoxC5 maps to a cluster of Hox proteins on chromosome 12 that are essential for morphogenesis. Encoding over 1,100 genes within 132 million bases, chromosome 12 makes up about 4.5% of the human genome. A number of skeletal deformities are linked to chromosome 12, including hypochondrogenesis, achondrogenesis and Kniest dysplasia.

REFERENCES

1. Acampora, D., et al. 1989. The human Hox gene family. *Nucleic Acids Res.* 17: 10385-10402.
2. McAlpine, P.J. and Shows, T.B. 1990. Nomenclature for human homeobox genes. *Genomics* 7: 460.
3. Arcioni, L., et al. 1992. The upstream region of the human homeobox gene HOX3D is a target for regulation by retinoic acid and Hox homeoproteins. *EMBO J.* 11: 265-277.
4. Guazzi, S., et al. 1994. The thyroid transcription factor-1 gene is a candidate target for regulation by Hox proteins. *EMBO J.* 13: 3339-3347.
5. Apiou, F., et al. 1996. Fine mapping of human Hox gene clusters. *Cytogenet. Cell Genet.* 73: 114-115.
6. Kosaki, K., et al. 2002. Complete mutation analysis panel of the 39 human Hox genes. *Teratology* 65: 50-62.
7. Online Mendelian Inheritance in Man, OMIM[™]. 2002. Johns Hopkins University, Baltimore, MD. MIM Number: 142973. World Wide Web URL: <http://www.ncbi.nlm.nih.gov/omim/>
8. Gong, L.G., et al. 2005. Analysis of single nucleotide polymorphisms and haplotypes in HoxC gene cluster within susceptible region 12q13 of simple congenital heart disease. *Zhonghua Yi Xue Yi Chuan Xue Za Zhi* 22: 497-501.

CHROMOSOMAL LOCATION

Genetic locus: HOXC5 (human) mapping to 12q13.13.

PRODUCT

HoxC5 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 HoxC5 shRNA Plasmid (h): sc-75287-SH and HoxC5 shRNA (h) Lentiviral Particles: sc-75287-V as alternate gene silencing products.

For independent verification of HoxC5 (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-75287A, sc-75287B and sc-75287C.

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

HoxC5 siRNA (h) is recommended for the inhibition of HoxC5 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

HoxC5 (1E10): sc-517171 is recommended as a control antibody for monitoring of HoxC5 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).

To ensure optimal results, the following support reagents are recommended: 1) Western Blotting: use m-IgG κ BP-HRP: sc-516102 or m-IgG κ BP-HRP (Cruz Marker): sc-516102-CM (dilution range: 1:1000-1:10000), Cruz Marker[™] Molecular Weight Standards: sc-2035, UltraCruz[®] Blocking Reagent: sc-516214 and Western Blotting Luminol Reagent: sc-2048. 2) Immunofluorescence: use m-IgG κ BP-FITC: sc-516140 or m-IgG κ BP-PE: sc-516141 (dilution range: 1:50-1:200) with UltraCruz[®] Mounting Medium: sc-24941 or UltraCruz[®] Hard-set Mounting Medium: sc-359850.

RT-PCR REAGENTS

Semi-quantitative RT-PCR may be performed to monitor HoxC5 gene expression knockdown using RT-PCR Primer: HoxC5 (h)-PR: sc-75287-PR (20 μ l, 515 bp). Annealing temperature for the primers should be 55-60° C and the extension temperature should be 68-72° C.

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.