

PLEKHM1 siRNA (m): sc-152320

BACKGROUND

PLEKHM1 (Pleckstrin homology domain-containing family M member 1), also known as 162 kDa adapter protein, is a 779 amino acid protein that contains two Pleckstrin homology (PH) domains, which are found in proteins that are involved in intracellular signaling. PLEKHM1 is located in the cytoplasm where it is involved in osteoclast vesicular transport and is therefore an essential protein for bone resorption. Individuals having defects in PLEKHM1 are afflicted with osteopetrosis autosomal recessive type 6 (OPTB6), a rare genetic disease that is characterized by abnormally dense bone due to ineffective bone resorption. Specifically, it is likely that mutations in the PLEKHM1 gene affect endosomal acidification/maturation and TRACP exocytosis, which has implications on osteoclast-osteoblast cross-talk.

REFERENCES

1. Hartel-Schenk, S., et al. 2001. Novel adapter protein AP162 connects a sialyl-Le_x-positive mucin with an apoptotic signal transduction pathway. *Glycoconj. J.* 18: 915-923.
2. Katoh, M., et al. 2004. Characterization of RUSC1 and RUSC2 genes in silico. *Oncol. Rep.* 12: 933-938.
3. Frattini, A., et al. 2007. The dissection of human autosomal recessive osteopetrosis identifies an osteoclast-poor form due to RANKL deficiency. *Cell Cycle* 6: 3027-3033.
4. Van Wesenbeeck, L., et al. 2007. Involvement of PLEKHM1 in osteoclastic vesicular transport and osteopetrosis in incisors absent rats and humans. *J. Clin. Invest.* 117: 919-930.
5. Online Mendelian Inheritance in Man, OMIM™. 2007. Johns Hopkins University, Baltimore, MD. MIM Number: 611466. World Wide Web URL: <http://www.ncbi.nlm.nih.gov/omim/>
6. Del Fattore, A., et al. 2008. Genetics, pathogenesis and complications of osteopetrosis. *Bone* 42: 19-29.
7. Del Fattore, A., et al. 2008. A new heterozygous mutation (R714C) of the osteopetrosis gene, Pleckstrin homolog domain containing family M (with run domain) member 1 (PLEKHM1), impairs vesicular acidification and increases TRACP secretion in osteoclasts. *J. Bone Miner. Res.* 23: 380-391.

CHROMOSOMAL LOCATION

Genetic locus: Plekhl1 (mouse) mapping to 11 E1.

PRODUCT

PLEKHM1 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 PLEKHM1 shRNA Plasmid (m): sc-152320-SH and PLEKHM1 shRNA (m) Lentiviral Particles: sc-152320-V as alternate gene silencing products.

For independent verification of PLEKHM1 (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-152320A, sc-152320B and sc-152320C.

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

PLEKHM1 siRNA (m) is recommended for the inhibition of PLEKHM1 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 PLEKHM1 gene expression knockdown using RT-PCR Primer: PLEKHM1 (m)-PR: sc-152320-PR (20 μ l). Annealing temperature for the primers should be 55-60° C and the extension temperature should be 68-72° C.

SELECT PRODUCT CITATIONS

1. Viereck, J., et al. 2016. Long noncoding RNA Chast promotes cardiac remodeling. *Sci. Transl. Med.* 8: 326ra22.

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.