

ATP13A1 siRNA (h): sc-97321

BACKGROUND

ATP13A1 (ATPase type 13A1), also known as probable cation-transporting ATPase 13A1 or ATP13A, is a 1,204 amino acid multi-pass membrane protein belonging to the cation transport ATPase (P-type) family and the type V sub-family. Highly conserved, ATP13A1 exists as three alternatively spliced isoforms. ATP13A1 participates in ATP, metal ion and nucleotide binding, ATPase activity, and hydrolase activity, such as acting on acid anhydrides and catalyzing transmembrane substance movement. ATP13A1 is encoded by a gene that maps to human chromosome 19p13.11. Chromosome 19 consists of approximately 63 million bases and makes up over 2% of human genomic DNA. Chromosome 19 contains the greatest gene density of the human chromosomes and is the genetic home for a number of immunoglobulin superfamily members, including killer cell and leukocyte Ig-like receptors, ICAMs, the CEACAM and PSG families, and Fc α receptors.

REFERENCES

- Schultheis, P.J., et al. 2004. Characterization of the P5 subfamily of P-type transport ATPases in mice. *Biochem. Biophys. Res. Commun.* 323: 731-738.
- Grimwood, J., et al. 2004. The DNA sequence and biology of human chromosome 19. *Nature* 428: 529-535.
- Kwasnicka-Crawford, D.A., et al. 2005. Characterization of a novel cation transporter ATPase gene (ATP13A4) interrupted by 3q25-q29 inversion in an individual with language delay. *Genomics* 86: 182-194.
- Wolfe, D.M., et al. 2006. Channeling studies in yeast: yeast as a model for channelopathies? *Neuromolecular Med.* 8: 279-306.
- Wenge, B., et al. 2008. Separation of membrane proteins by two-dimensional electrophoresis using cationic rehydrated strips. *Electrophoresis* 29: 1511-1517.
- Nath, A.K., et al. 2009. Proteomic-based detection of a protein cluster dysregulated during cardiovascular development identifies biomarkers of congenital heart defects. *PLoS ONE* 4: e4221.
- Chadi, S., et al. 2010. Brain transcriptional stability upon prion protein-encoding gene invalidation in zygotic or adult mouse. *BMC Genomics* 11: 448.
- Vallipuram, J., et al. 2010. The E646D-ATP13A4 mutation associated with autism reveals a defect in calcium regulation. *Cell. Mol. Neurobiol.* 30: 233-246.
- Landi, D., et al. 2011. Prediction of the biological effect of polymorphisms within microRNA binding sites. *Methods Mol. Biol.* 676: 197-210.

CHROMOSOMAL LOCATION

Genetic locus: ATP13A1 (human) mapping to 19p13.11.

PROTOCOLS

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

PRODUCT

ATP13A1 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 ATP13A1 shRNA Plasmid (h): sc-97321-SH and ATP13A1 shRNA (h) Lentiviral Particles: sc-97321-V as alternate gene silencing products.

For independent verification of ATP13A1 (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-97321A, sc-97321B and sc-97321C.

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

ATP13A1 siRNA (h) is recommended for the inhibition of ATP13A1 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.

RT-PCR REAGENTS

Semi-quantitative RT-PCR may be performed to monitor ATP13A1 gene expression knockdown using RT-PCR Primer: ATP13A1 (h)-PR: sc-97321-PR (20 μ l). 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.