

# DIP2A siRNA (m): sc-62213

## BACKGROUND

DIP2A (disco-interacting protein 2 homolog A), also known as DIP2, is a 1571 amino acid nuclear protein. It is one of three human homologs (DIP2A, DIP2B and DIP2C) of the *Drosophila* dip2 (disconnected-interacting protein 2) protein. In *Drosophila*, dip2 interacts with disco, a protein required for neuronal connections in the visual systems of larvae and adults. The closest vertebrate homologs to disco are the basophilin genes. In mice, DIP2 homologs show restricted expression to the brain. This suggests that, similar to the function of *Drosophila* dip2, vertebrate DIP2 homologs may play a role in the development of the nervous system. Expressed ubiquitously with highest expression in the brain, DIP2A is thought to function in signaling throughout the central nervous system by providing positional clues for axon patterning and pathfinding. Four isoforms of DIP2A exist due to alternative splicing events.

## REFERENCES

1. Mukhopadhyay, M., et al. 2002. Cloning, genomic organization and expression pattern of a novel *Drosophila* gene, the disco-interacting protein 2 (dip2), and its murine homolog. *Gene* 293: 59-65.
2. Online Mendelian Inheritance in Man, OMIM<sup>™</sup>. 2002. Johns Hopkins University, Baltimore, MD. MIM Number: 607711. World Wide Web URL: <http://www.ncbi.nlm.nih.gov/omim/>
3. DeSousa, D., et al. 2003. A novel double-stranded RNA-binding protein, disco interacting protein 1 (DIP1), contributes to cell fate decisions during *Drosophila* development. *J. Biol. Chem.* 278: 38040-38050.
4. De Felice, B., et al. 2003. Characterization of DIP1, a novel nuclear protein in *Drosophila melanogaster*. *Biochem. Biophys. Res. Commun.* 307: 224-228.
5. Bondos, S.E., et al. 2004. Hox transcription factor ultrabithorax I $\beta$  physically and genetically interacts with disconnected interacting protein 1, a double-stranded RNA-binding protein. *J. Biol. Chem.* 279: 26433-26444.

## CHROMOSOMAL LOCATION

Genetic locus: Dip2a (mouse) mapping to 10 C1.

## PRODUCT

DIP2A 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  $\mu$ M solution once resuspended using protocol below. Suitable for 50-100 transfections. Also see DIP2A shRNA Plasmid (m): sc-62213-SH and DIP2A shRNA (m) Lentiviral Particles: sc-62213-V as alternate gene silencing products.

For independent verification of DIP2A (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-62213A, sc-62213B and sc-62213C.

## PROTOCOLS

See our web site at [www.scbt.com](http://www.scbt.com) or our catalog 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  $\mu$ l of the RNase-free water provided. Resuspension of the siRNA duplex in 330  $\mu$ l of RNase-free water makes a 10  $\mu$ M solution in a 10  $\mu$ M Tris-HCl, pH 8.0, 20 mM NaCl, 1 mM EDTA buffered solution.

## APPLICATIONS

DIP2A siRNA (m) is recommended for the inhibition of DIP2A 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  $\mu$ M in 66  $\mu$ 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 DIP2A gene expression knockdown using RT-PCR Primer: DIP2A (m)-PR: sc-62213-PR (20  $\mu$ 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.