

## **Product datasheet for SR317279**

## OriGene Technologies, Inc.

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## **CNPY1 Human siRNA Oligo Duplex (Locus ID 285888)**

**Product data:** 

**Product Type:** siRNA Oligo Duplexes

Purity: HPLC purified

**Quality Control:** Tested by ESI-MS

Sequences: Available with shipment

**Stability:** One year from date of shipment when stored at -20°C.

# of transfections: Approximately 330 transfections/2nmol in 24-well plate under optimized conditions (final

conc. 10 nM).

**Note:** Single siRNA duplex (10nmol) can be ordered.

**RefSeq:** <u>NM 001103176, NM 001369814, NR 163164, NM 001369813, NM 001369815,</u>

NM 001369816, NM 001369820, NR 163159, NR 163161, NR 163162, NR 163165, NR 163167, NM 001369818, NM 001369821, NR 163157, NR 163158, NR 163160,

NR 163163, NR 163166

UniProt ID: Q3B712

Components: CNPY1 (Human) - 3 unique 27mer siRNA duplexes - 2 nmol each (Locus ID 285888)

Included - SR30004, Trilencer-27 Universal Scrambled Negative Control siRNA Duplex - 2 nmol

Included - SR30005, RNAse free siRNA Duplex Resuspension Buffer - 2 ml

Summary: Cnpy1 is expressed in the midbrain-hindbrain (MHB) boundary in zebrafish, binds FGFR1

(MIM 136350), and plays a role in FGF signaling (Hirate and Okamoto, 2006 [PubMed

16488878]).[supplied by OMIM, Dec 2008]



## Performance Guaranteed:

OriGene guarantees that at least two of the three Dicer-Substrate duplexes in the kit will provide at least 70% or more knockdown of the target mRNA when used at 10 nM concentration by quantitative RT-PCR when the TYE-563 fluorescent transfection control duplex (cat# SR30002) indicates that >90% of the cells have been transfected and the HPRT positive control (cat# SR30003) provides 90% knockdown efficiency.

For non-conforming siRNA, requests for replacement product must be made within ninety (90) days from the date of delivery of the siRNA kit. To arrange for a free replacement with newly designed duplexes, please contact Technical Services at techsupport@origene.com. Please provide your data indicating the transfection efficiency and measurement of gene expression knockdown compared to the scrambled siRNA control (quantitative RT-PCR data required).