

## **Product datasheet for SR412577**

## OriGene Technologies, Inc.

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## **Cep41 Mouse siRNA Oligo Duplex (Locus ID 83922)**

**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:** NM 031998, NR 130775, NR 130781

UniProt ID: Q99NF3

**Synonyms:** 1700017E11Rik; 2810431D15Rik; Cep4; Tsga; Tsga14

Components: Cep41 (Mouse) - 3 unique 27mer siRNA duplexes - 2 nmol each (Locus ID 83922)

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

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

**Summary:** This gene encodes a centrosomal protein which may be required for tubulin glutamylation in

cilia during ciliogenesis. Mutations in a similar gene in human have been associated with

Joubert Syndrome 15, an autosomal recessive ciliopathy and neurological disorder.

Alternative splicing results in multiple transcript variants. [provided by RefSeq, Feb 2015]

**Performance** OriGene guarantees that at least two of the three Dicer-Substrate duplexes in the kit will

**Guaranteed:** 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).

