

# **Product datasheet for SR317270**

## OriGene Technologies, Inc.

9620 Medical Center Drive, Ste 200 Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com CN: techsupport@origene.cn

## C6ORF182 (CEP57L1) Human siRNA Oligo Duplex (Locus ID 285753)

#### **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 001083535, NM 001271852, NM 001271853, NM 173830, NM 001350652,

NM 001350653, NM 001350654, NM 001350655, NM 001350656, NM 001350657, NM 001350658, NM 001350659, NM 001350660, NM 001350661, NM 001350662,

NM 001350663, NM 001350664, NM 001350665, NM 001350666, NR 146888, NR 146889

UniProt ID: Q8IYX8

Synonyms: bA487F23.2; C6orf182; cep57R

Components: CEP57L1 (Human) - 3 unique 27mer siRNA duplexes - 2 nmol each (Locus ID 285753)

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

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

**Summary:** Centrosomal protein which may be required for microtubule attachment to centrosomes.

[UniProtKB/Swiss-Prot Function]





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# 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).