

Product datasheet for SR314581

OriGene Technologies, Inc.

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C1orf19 (TSEN15) Human siRNA Oligo Duplex (Locus ID 116461)

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 001127394, NM 001300764, NM 001300766, NM 052965, NR 023349, NR 125335,

NM 001363643

UniProt ID: Q8WW01

Synonyms: C1orf19; PCH2F; sen15

Components: TSEN15 (Human) - 3 unique 27mer siRNA duplexes - 2 nmol each (Locus ID 116461)

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 subunit of the tRNA splicing endonuclease, which catalyzes the removal

of introns from tRNA precursors. Alternative splicing results in multiple transcript variants. There is a pseudogene of this gene on chromosome 17. [provided by RefSeq, Jul 2014]

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

or order at least 70% of more knockdown of the target mixth when ased at 10 mix

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).

