

## **Product datasheet for SR304667**

## OriGene Technologies, Inc.

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## **SURF5 (MED22) Human siRNA Oligo Duplex (Locus ID 6837)**

**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 006752</u>, <u>NM 133640</u>, <u>NM 181491</u>

UniProt ID: Q15528

Synonyms: MED24, MGC48682

Components: MED22 (Human) - 3 unique 27mer siRNA duplexes - 2 nmol each (Locus ID 6837)

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 protein component of the mediator complex, which functions in the

regulation of transcription by bridging interactions between gene-specific regulatory factors, RNA polymerase II, and general transcription factors. Alternatively spliced transcript variants

encoding different isoforms have been observed. [provided by RefSeq, Jul 2013]

**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).

