

Product datasheet for SR311791

OriGene Technologies, Inc.

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TLNRD1 Human siRNA Oligo Duplex (Locus ID 59274)

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_022566</u>

UniProt ID: Q9H1K6

Synonyms: MESDC1

Components: MESDC1 (Human) - 3 unique 27mer siRNA duplexes - 2 nmol each (Locus ID 59274)

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 that is regulated by micro RNA MiR-574-3, and is thought to have

an oncogenic function in human bladder cancer. A similar gene in mouse is located in a chromosomal region critical for differentiation of mesoderm, which affects embryo patterning and the formation of heart, muscle, blood, skeleton and the urogenital system. The mouse gene is expressed in early development, and in the adult. [provided by RefSeq, Nov 2016]



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).