

## Product datasheet for SR312159

## OriGene Technologies, Inc.

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## DCLRE1B Human siRNA Oligo Duplex (Locus ID 64858)

## **Product data:**

**Product Type:** siRNA Oligo Duplexes

**HPLC** purified **Purity:** 

**Quality Control:** Tested by ESI-MS

Available with shipment **Sequences:** 

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).

Single siRNA duplex (10nmol) can be ordered. Note:

RefSeq: NM 001319946, NM 001319947, NM 022836, NM 001363690, NM 001363691

**UniProt ID:** Q9H816

Synonyms: APOLLO; SNM1B; SNMIB

Components: DCLRE1B (Human) - 3 unique 27mer siRNA duplexes - 2 nmol each (Locus ID 64858)

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

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

DNA interstrand cross-links prevent strand separation, thereby physically blocking **Summary:** 

> transcription, replication, and segregation of DNA. DCLRE1B is one of several evolutionarily conserved genes involved in repair of interstrand cross-links (Dronkert et al., 2000 [PubMed

10848582]).[supplied by OMIM, Mar 2008]

**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).

