

## **Product datasheet for SR423617**

## OriGene Technologies, Inc.

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## **Bod1l Mouse siRNA Oligo Duplex (Locus ID 665775)**

**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\_001081422</u>, <u>NM\_001310601</u>

UniProt ID: E9Q6J5

**Synonyms:** A230054D04Rik; AI853319; FAM44A; mKIAA1327

Components: Bod1l (Mouse) - 3 unique 27mer siRNA duplexes - 2 nmol each (Locus ID 665775)

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

replication forks and protecting RAD51 nucleofilaments from the antirecombinogenic activities

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

**Summary:** Component of the fork protection machinery required to protect stalled/damaged replication

forks from uncontrolled DNA2-dependent resection. Acts by stabilizing RAD51 at stalled

of FBHI and BLM. Does not regulate spindle orientation.[UniProtKB/Swiss-Prot Function]





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).