

## **Product datasheet for SR416311**

### OriGene Technologies, Inc.

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## Midn Mouse siRNA Oligo Duplex (Locus ID 59090)

#### **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 021565</u>

UniProt ID: Q3TPJ7

**Synonyms:** 3000003C15Rik

Components: Midn (Mouse) - 3 unique 27mer siRNA duplexes - 2 nmol each (Locus ID 59090)

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 contains an ubiquitin-like domain. This protein may be

involved in the regulation of brain development as inferred by its high expression level in the embryonic midbrain. This protein has been found to negatively regulate glucokinase activity and insulin secretion in pancreatic beta cells. Alternative splicing results in multiple transcript

variants. [provided by RefSeq, Apr 2015]







# 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).