

## **Product datasheet for SR307097**

## OriGene Technologies, Inc.

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## MAD2L2 Human siRNA Oligo Duplex (Locus ID 10459)

## **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 001127325</u>, <u>NM 006341</u>

UniProt ID: Q9UI95

**Synonyms:** FANCV; MAD2B; POLZ2; REV7

Components: MAD2L2 (Human) - 3 unique 27mer siRNA duplexes - 2 nmol each (Locus ID 10459)

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

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

**Summary:** The protein encoded by this gene is a component of the mitotic spindle assembly checkpoint

that prevents the onset of anaphase until all chromosomes are properly aligned at the metaphase plate. The encoded protein, which is similar to MAD2L1, is capable of interacting

with ADAM9, ADAM15, REV1, and REV3 proteins. [provided by RefSeq, Jul 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).

