

## Product datasheet for SR314207

## OriGene Technologies, Inc.

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## MOB4A (MOB1B) Human siRNA Oligo Duplex (Locus ID 92597)

## **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 001244766, NM 001244767, NM 173468

**UniProt ID:** Q7L9L4

Synonyms: MATS2; MOB4A; MOBKL1A

Components: MOB1B (Human) - 3 unique 27mer siRNA duplexes - 2 nmol each (Locus ID 92597)

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

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

The protein encoded by this gene is similar to the yeast Mob1 protein. Yeast Mob1 binds **Summary:** 

> Mps1p, a protein kinase essential for spindle pole body duplication and mitotic checkpoint regulation. Three transcript variants encoding different isoforms have been found for this

gene. [provided by RefSeq, Sep 2011]

**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).

