

## **Product datasheet for SR302772**

## OriGene Technologies, Inc.

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## SMAD9 Human siRNA Oligo Duplex (Locus ID 4093)

**Product data:** 

**Guaranteed:** 

**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 001127217</u>, <u>NM 005905</u>

**UniProt ID:** <u>O15198</u>

Synonyms: MADH6; MADH9; PPH2; SMAD8; SMAD8/9; SMAD8A; SMAD8B

Components: SMAD9 (Human) - 3 unique 27mer siRNA duplexes - 2 nmol each (Locus ID 4093)

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 member of the SMAD family, which transduces signals

from TGF-beta family members. The encoded protein is activated by bone morphogenetic proteins and interacts with SMAD4. Two transcript variants encoding different isoforms have

been found for this gene.[provided by RefSeq, Jan 2010]

**Performance** 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).

