

## Product datasheet for SR413891

## OriGene Technologies, Inc.

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## Elmo1 Mouse siRNA Oligo Duplex (Locus ID 140580)

## **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 080288, NM 198093, NR 038122

**UniProt ID:** Q8BPU7

Synonyms: 6330578D22Rik; C230095H21Rik; CED-12

Components: Elmo1 (Mouse) - 3 unique 27mer siRNA duplexes - 2 nmol each (Locus ID 140580)

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

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

Involved in cytoskeletal rearrangements required for phagocytosis of apoptotic cells and cell **Summary:** 

> motility. Acts in association with DOCK1 and CRK. Was initially proposed to be required in complex with DOCK1 to activate Rac Rho small GTPases. May enhance the guanine nucleotide

exchange factor (GEF) activity of DOCK1 (By similarity).[UniProtKB/Swiss-Prot Function]

**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).

