

Product datasheet for SR513786

OriGene Technologies, Inc.

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Panx1 Rat siRNA Oligo Duplex (Locus ID 315435)

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:
 NM 001270548, NM 001270549, NM 199397

UniProt ID: <u>P60570</u>

Synonyms: px1

Components: Panx1 (Rat) - 3 unique 27mer siRNA duplexes - 2 nmol each (Locus ID 315435)

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 plasma membrane protein that is a structural

component of gap junctions. The encoded protein acts as a homodimer or as a heterodimer with other isoforms or proteins. Two additional variants have been found, and the isoforms expressed from them are found in the cytoplasm. It is thought that these two isoforms could

attenuate the actions of the membrane-bound protein. [provided by RefSeq, Jul 2012]





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).