

Product datasheet for SR427122

OriGene Technologies, Inc.

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Lrmp Mouse siRNA Oligo Duplex (Locus ID 16970)

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: NM 001281980, NM 001281981, NM 008511, NM 001361633, NM 001368864

UniProt ID: Q60664

Synonyms: D6Int3; D6Int4; D6Int5; D6Int7; D6Int8; Jaw1

Components: Lrmp (Mouse) - 3 unique 27mer siRNA duplexes - 2 nmol each (Locus ID 16970)

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

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

Summary: Plays a role in the delivery of peptides to major histocompatibility complex (MHC) class I

molecules; this occurs in a transporter associated with antigen processing (TAP)-independent

manner. May play a role in taste signal transduction via ITPR3. May play a role during fertilization in pronucleus congression and fusion.[UniProtKB/Swiss-Prot Function]

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).

