

Product datasheet for SR422721

OriGene Technologies, Inc.

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Mtus2 Mouse siRNA Oligo Duplex (Locus ID 77521)

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 001033601, NM 029920, NM 001359503, NM 001359505, NM 001359511

UniProt ID: Q3UHD3

Synonyms: Gm763, AU067634, mKIAA0774, 5730592G18Rik, A730013O20Rik

Components: Mtus2 (Mouse) - 3 unique 27mer siRNA duplexes - 2 nmol each (Locus ID 77521)

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

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

Summary: Binds microtubules. Together with MAPRE1 may target the microtubule depolymerase KIF2C

to the plus-end of microtubules. May regulate the dynamics of microtubules at their growing

to the plus-end of file occupates. May regulate the dynamics of file occupates at their growing

distal tip (By similarity).[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).

