

Product datasheet for SR421014

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Ttll7 Mouse siRNA Oligo Duplex (Locus ID 70892)

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 001302957, NM 001302958, NM 027594, NM 001357217

UniProt ID: A4Q9F0

Synonyms: 1110049N09Rik; 4921517B04Rik; C630030B20; mTTLL7

Components: Ttll7 (Mouse) - 3 unique 27mer siRNA duplexes - 2 nmol each (Locus ID 70892)

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

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

Summary: Polyglutamylase which preferentially modifies beta-tubulin (PubMed:16901895,

PubMed:17499049, PubMed:19152315). Mediates both ATP-dependent initiation and elongation of polyglutamylation of microtubules (PubMed:16901895, PubMed:19152315). Required for neurite growth; responsible for the strong increase in tubulin polyglutamylation during postnatal neuronal maturation (PubMed:16901895).[UniProtKB/Swiss-Prot Function]





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).