## Product datasheet for SR417758

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## Has1 Mouse siRNA Oligo Duplex (Locus ID 15116)

## Product data:

Product Type:
Purity:
Quality Control:
Sequences:
Stability:
\# of transfections:

Note:
RefSeq:
UniProt ID:
Synonyms:
Components:

Summary:
siRNA Oligo Duplexes
HPLC purified
Tested by ESI-MS
Available with shipment
One year from date of shipment when stored at $-20^{\circ} \mathrm{C}$.
Approximately 330 transfections/2nmol in 24 -well plate under optimized conditions (final conc. 10 nM ).
Single siRNA duplex (10nmol) can be ordered.
NM 008215
Q61647
HAS
Has1 (Mouse) - 3 unique 27mer siRNA duplexes - 2 nmol each (Locus ID 15116) Included - SR30004, Trilencer-27 Universal Scrambled Negative Control siRNA Duplex - 2 nmol Included - SR30005, RNAse free siRNA Duplex Resuspension Buffer - 2 ml

Catalyzes the addition of GlcNAc or GlcUA monosaccharides to the nascent hyaluronan polymer. Therefore, it is essential to hyaluronan synthesis a major component of most extracellular matrices that has a structural role in tissues architectures and regulates cell adhesion, migration and differentiation. This is one of the isozymes catalyzing that reaction. Also able to catalyze the synthesis of chito-oligosaccharide depending on the substrate. [UniProtKB/Swiss-Prot Function]

## Performance <br> 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).

