

Product datasheet for SR422928

Abcc6 Mouse siRNA Oligo Duplex (Locus ID 27421)

Product data:

OriGene Technologies, Inc.

9620 Medical Center Drive, Ste 200 Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com CN: techsupport@origene.cn

siRNA Oligo Duplexes
HPLC purified
Tested by ESI-MS
Available with shipment
One year from date of shipment when stored at -20°C.
Approximately 330 transfections/2nmol in 24-well plate under optimized conditions (final conc. 10 nM).
Single siRNA duplex (10nmol) can be ordered.
<u>NM 018795</u>
<u>Q9R1S7</u>
Abcc1b; DCC; Dysca; dyscalc; Dyscalc1; Mr; Mrp6
Abcc6 (Mouse) - 3 unique 27mer siRNA duplexes - 2 nmol each (Locus ID 27421) Included - SR30004, Trilencer-27 Universal Scrambled Negative Control siRNA Duplex - 2 nmol Included - SR30005, RNAse free siRNA Duplex Resuspension Buffer - 2 ml
The protein encoded by this gene is a member of the superfamily of ATP-binding cassette (ABC) transporters. ABC proteins transport various molecules across extra- and intra-cellular membranes. ABC genes are divided into seven distinct subfamilies (ABC1, MDR/TAP, MRP, ALD, OABP, GCN20, White). This protein is a member of the MRP subfamily which is involved in multi-drug resistance. The specific function of this protein is unknown; however, a similar rat protein has been identified as the major canalicular bile salt export pump of liver. [provided by RefSeq, Jul 2008]



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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).

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