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Product datasheet for SR417590

Ahrr Mouse siRNA Oligo Duplex (Locus ID 11624)

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_009644
UniProt ID:	Q3U1U7
Synonyms:	mKIAA1234
Components:	Ahrr (Mouse) - 3 unique 27mer siRNA duplexes - 2 nmol each (Locus ID 11624) Included - SR30004, Trilencer-27 Universal Scrambled Negative Control siRNA Duplex - 2 nmol Included - SR30005, RNAse free siRNA Duplex Resuspension Buffer - 2 ml
Summary:	This gene encodes a protein that represses aryl hydrocarbon receptor-dependent signaling. The encoded protein competes with the aryl hydrocarbon receptor transcription factor for heterodimerization with the aryl hydrocarbon receptor nuclear translocator protein and binding to xenobiotic response element (XRE) sequence in many genes. This protein is implicated in the regulation of cell growth and differentiation as well as mediating dioxin toxicity. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Nov 2015]

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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).