

# Product datasheet for SR416659

## Adgrg5 Mouse siRNA Oligo Duplex (Locus ID 382045)

### **Product data:**

#### **Product Type:** siRNA Oligo Duplexes HPLC purified **Purity: Quality Control:** Tested by ESI-MS Available with shipment Sequences: 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. NM 001033468, NM 001145972 **RefSeq: UniProt ID:** Q3V3Z3 Synonyms: Gm1109; Gpr114; PGR27 **Components:** Adgrg5 (Mouse) - 3 unique 27mer siRNA duplexes - 2 nmol each (Locus ID 382045) Included - SR30004, Trilencer-27 Universal Scrambled Negative Control siRNA Duplex - 2 nmol Included - SR30005, RNAse free siRNA Duplex Resuspension Buffer - 2 ml Adhesion G protein-coupled receptor (GPCR). Transduces intracellular signals through Summary: coupling to guanine nucleotide-binding protein G(s) subunit alpha and activation of adenylate cyclase pathway (PubMed:22575658). Isoform 1, but not isoform 2, is constitutively active, as evidenced by elevated basal cAMP levels, and responds to mechanical activation (shaking) (PubMed:26499266).[UniProtKB/Swiss-Prot Function]



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#### OriGene Technologies, Inc.

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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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