

Product datasheet for SR300396

OriGene Technologies, Inc.

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BAI2 (ADGRB2) Human siRNA Oligo Duplex (Locus ID 576)

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: <u>NM 001294335</u>, <u>NM 001294336</u>, <u>NM 001703</u>, <u>NM 001364857</u>

UniProt ID: O60241
Synonyms: BAI2

Components: ADGRB2 (Human) - 3 unique 27mer siRNA duplexes - 2 nmol each (Locus ID 576)

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 a seven-span transmembrane protein that is thought to be a member of

the secretin receptor family. The encoded protein is a brain-specific inhibitor of angiogenesis.

The mature peptide may be further cleaved into additional products (PMID:20367554). Alternative splicing results in multiple transcript variants. [provided by RefSeq, Jun 2014]

Performance OriGene guarantees that at least two of the three Dicer-Substrate duplexes in the kit will

Guaranteed: 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).

