

Product datasheet for SR303259

OriGene Technologies, Inc.

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Nucleobindin 2 (NUCB2) Human siRNA Oligo Duplex (Locus ID 4925)

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 001330227, NM 005013, NM 001352661, NM 001352662, NM 001352663,

NM 001352664, NM 001352665, NM 001352666, NM 001352667, NM 001352668,

NM 001352669, NM 001352670, NM 001352671, NM 001352672

UniProt ID: P80303

Synonyms: HEL-S-109; NEFA

Components: NUCB2 (Human) - 3 unique 27mer siRNA duplexes - 2 nmol each (Locus ID 4925)

Included - SR30004, Trilencer-27 Universal Scrambled Negative Control siRNA Duplex - 2 nmol

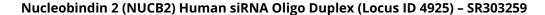
Included - SR30005, RNAse free siRNA Duplex Resuspension Buffer - 2 ml

Summary: This gene encodes a protein with a suggested role in calcium level maintenance, eating

regulation in the hypothalamus, and release of tumor necrosis factor from vascular endothelial cells. This protein binds calcium and has EF-folding domains. [provided by

RefSeq, Oct 2011]







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).