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Product datasheet for SR301418

EPB4IL2 (EPB41L2) Human siRNA Oligo Duplex (Locus ID 2037)

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_001135554, NM_001135555, NM_001199388, NM_001199389, NM_001252660, NM_001431, NM_001350299, NM_001350301, NM_001350302, NM_001350303, NM_001350304, NM_001350305, NM_001350306, NM_001350307, NM_001350308, NM_001350309, NM_001350310, NM_001350311, NM_001350312, NM_001350313, NM_001350314, NM_001350315, NM_001350320, NR_146620
UniProt ID:	O43491
Synonyms:	4.1-G; 4.1G
Components:	EPB41L2 (Human) - 3 unique 27mer siRNA duplexes - 2 nmol each (Locus ID 2037) Included - SR30004, Trilencer-27 Universal Scrambled Negative Control siRNA Duplex - 2 nmol Included - SR30005, RNAse free siRNA Duplex Resuspension Buffer - 2 ml
Summary:	Required for dynein-dynactin complex and NUMA1 recruitment at the mitotic cell cortex during anaphase (PubMed:23870127).[UniProtKB/Swiss-Prot Function]

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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).