

Product datasheet for SR311448

OriGene Technologies, Inc.

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ZNF304 Human siRNA Oligo Duplex (Locus ID 57343)

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_001290318, NM_001290319, NM_020657, NM_001329456</u>

UniProt ID: Q9HCX3

Components: ZNF304 (Human) - 3 unique 27mer siRNA duplexes - 2 nmol each (Locus ID 57343)

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 member of the Krueppel C2H2-type zinc-finger family of proteins. The

encoded protein functions as a transcriptional repressor that recruits a corepressor complex

to stimulate promoter hypermethylation and transcriptional silencing of target genes.

Expression of this gene is upregulated in colorectal, ovarian and breast cancer, and this gene

may promote cancer cell survival, growth and invasion. [provided by RefSeq, Jul 2016]



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).