

# Product datasheet for SR305949

## CLIC3 Human siRNA Oligo Duplex (Locus ID 9022)

### **Product data:**

### OriGene Technologies, Inc.

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siRNA Oligo Duplexes
HPLC purified
Tested by ESI-MS
Available with shipment
One year from date of shipment when stored at -20°C.
Approximately 330 transfections/2nmol in 24-well plate under optimized conditions (final conc. 10 nM).
Single siRNA duplex (10nmol) can be ordered.
<u>NM 004669</u>
<u>O95833</u>
CLIC3 (Human) - 3 unique 27mer siRNA duplexes - 2 nmol each (Locus ID 9022) Included - SR30004, Trilencer-27 Universal Scrambled Negative Control siRNA Duplex - 2 nmol Included - SR30005, RNAse free siRNA Duplex Resuspension Buffer - 2 ml
Chloride channels are a diverse group of proteins that regulate fundamental cellular processes including stabilization of cell membrane potential, transepithelial transport, maintenance of intracellular pH, and regulation of cell volume. Chloride intracellular channel 3 is a member of the p64 family and is predominantly localized in the nucleus and stimulates chloride ion channel activity. In addition, this protein may participate in cellular growth control, based on its association with ERK7, a member of the MAP kinase family. [provided by RefSeq, Jul 2008]



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# CLIC3 Human siRNA Oligo Duplex (Locus ID 9022) - SR305949Performance<br/>Guaranteed:OriGene guarantees that at least two of the three Dicer-Substrate duplexes in the kit will<br/>provide at least 70% or more knockdown of the target mRNA when used at 10 nM<br/>concentration by quantitative RT-PCR when the TYE-563 fluorescent transfection control<br/>duplex (cat# SR30002) indicates that >90% of the cells have been transfected and the HPRT<br/>positive control (cat# SR30003) provides 90% knockdown efficiency.For non-conforming siRNA, requests for replacement product must be made within ninety<br/>(90) days from the date of delivery of the siRNA kit. To arrange for a free replacement with<br/>newly designed duplexes, please contact Technical Services at techsupport@origene.com.<br/>Please provide your data indicating the transfection efficiency and measurement of gene<br/>expression knockdown compared to the scrambled siRNA control (quantitative RT-PCR data

required).

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