

### **Product datasheet for SR303933**

## OriGene Technologies, Inc.

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# PCYT2 Human siRNA Oligo Duplex (Locus ID 5833)

#### **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 001184917, NM 001256433, NM 001256434, NM 001256435, NM 001282203,

NM 001282204, NM 001330518, NM 002861, NR 033681, NR 033682, NR 033683,

NR 033685

UniProt ID: Q99447
Synonyms: ET; SPG82

Components: PCYT2 (Human) - 3 unique 27mer siRNA duplexes - 2 nmol each (Locus ID 5833)

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 an enzyme that catalyzes the formation of CDP-ethanolamine from CTP

and phosphoethanolamine in the Kennedy pathway of phospholipid synthesis. Alternative

splicing results in multiple transcript variants. [provided by RefSeq, May 2010]





# 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).