

Product datasheet for **SR311285**

PARP6 Human siRNA Oligo Duplex (Locus ID 56965)

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_001323515 , NM_001323516 , NM_001323519 , NM_001323521 , NM_001323522 , NM_001323523 , NM_001323524 , NM_001323525 , NM_001323526 , NM_001323528 , NM_001323530 , NM_001323531 , NM_001323532 , NM_020213 , NM_020214 , NR_136594 , NR_136596 , NR_136599 , NR_136603 , NR_136604 , NR_136605 , NR_136606 , NR_136607 , NR_136608 , NR_136609 , NR_136610 , NR_136611
UniProt ID:	Q2NL67
Synonyms:	ARTD17; PARP-6-B1; PARP-6-C; pART17
Components:	PARP6 (Human) - 3 unique 27mer siRNA duplexes - 2 nmol each (Locus ID 56965) Included - SR30004, Trilencer-27 Universal Scrambled Negative Control siRNA Duplex - 2 nmol Included - SR30005, RNase free siRNA Duplex Resuspension Buffer - 2 ml
Summary:	Mono-ADP-ribosyltransferase that mediates mono-ADP-ribosylation of target proteins. [UniProtKB/Swiss-Prot Function]



[View online »](#)

**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).