

## Product datasheet for **SR322818**

### TP53I11 Human siRNA Oligo Duplex (Locus ID 9537)

#### 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:	<a href="#">NM_001076787</a> , <a href="#">NM_001258320</a> , <a href="#">NM_001258321</a> , <a href="#">NM_001258322</a> , <a href="#">NM_001258323</a> , <a href="#">NM_001258324</a> , <a href="#">NM_001318384</a> , <a href="#">NM_001318385</a> , <a href="#">NM_001318386</a> , <a href="#">NM_001318387</a> , <a href="#">NM_001318388</a> , <a href="#">NM_001318389</a> , <a href="#">NM_001318390</a> , <a href="#">NM_006034</a> , <a href="#">NR_134612</a>
UniProt ID:	<a href="#">Q14683</a>
Synonyms:	PIG11
Components:	TP53I11 (Human) - 3 unique 27mer siRNA duplexes - 2 nmol each (Locus ID 9537) Included - SR30004, Trilencer-27 Universal Scrambled Negative Control siRNA Duplex - 2 nmol Included - SR30005, RNAse free siRNA Duplex Resuspension Buffer - 2 ml
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](mailto: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).



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