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Product datasheet for TL312132V

INSL5 Human shRNA Lentiviral Particle (Locus ID 10022)

Product data:

Product Type:	shRNA Lentiviral Particles
Product Name:	INSL5 Human shRNA Lentiviral Particle (Locus ID 10022)
Locus ID:	10022
Synonyms:	PRO182; UNQ156
Vector:	pGFP-C-shLenti (TR30023)
Format:	Lentiviral particles
Components:	INSL5 - Human shRNA lentiviral particles (4 unique 29mer target-specific shRNA, 1 scramble control), 0.5 ml each, >10^7 TU/ml.
RefSeq:	NM_005478, NM_005478.1, NM_005478.2, NM_005478.3, NM_005478.4, BC101646, BC101646.1, BC101648
UniProt ID:	Q9Y5Q6
Summary:	The protein encoded by this gene contains a classical signature of the insulin superfamily and is highly similar to relaxin 3 (RLN3/INSL7). [provided by RefSeq, Jul 2008]
shRNA Design:	These shRNA constructs were designed against multiple splice variants at this gene locus. To be certain that your variant of interest is targeted, please contact techsupport@origene.com. If you need a special design or shRNA sequence, please utilize our custom shRNA service.
Performance Guaranteed:	OriGene guarantees that the sequences in the shRNA expression cassettes are verified to correspond to the target gene with 100% identity. One of the four constructs at minimum are guaranteed to produce 70% or more gene expression knock-down provided a minimum transfection efficiency of 80% is achieved. Western Blot data is recommended over qPCR to evaluate the silencing effect of the shRNA constructs 72 hrs post transfection. To properly assess knockdown, the gene expression level from the included scramble control vector must be used in comparison with the target-specific shRNA transfected samples. For non-conforming shRNA, requests for replacement product must be made within ninety (90) days from the date of delivery of the shRNA kit. To arrange for a free replacement with newly designed constructs, 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 shRNA control (Western Blot data preferred).

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