

Product datasheet for **TL320026V**

GAS8-AS1 Human shRNA Lentiviral Particle (Locus ID 750)

Product data:

Product Type:	shRNA Lentiviral Particles
Product Name:	GAS8-AS1 Human shRNA Lentiviral Particle (Locus ID 750)
Locus ID:	750
Synonyms:	MGC132509
Vector:	pGFP-C-shLenti (TR30023)
Format:	Lentiviral particles
Components:	C16orf3 - Human shRNA lentiviral particles (4 unique 29mer target-specific shRNA, 1 scramble control), 0.5 ml each, >10 ⁷ TU/ml.
RefSeq:	NM_001214 , NR_122031 , NM_001214.1 , NM_001214.2 , NM_001214.3 , BC093751 , BC104849
Summary:	This gene encodes a long non-coding RNA (lncRNA) that may function as a tumor suppressor. Mutations in this gene have been identified in human papillary thyroid carcinoma (PTC) patients that abrogate the ability of encoded lncRNA to inhibit cancer cell growth. [provided by RefSeq, May 2017]
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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