

Product datasheet for TL316948V

OriGene Technologies, Inc.

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SMAP (C11orf58) Human shRNA Lentiviral Particle (Locus ID 10944)

Product data:

Product Type: shRNA Lentiviral Particles

Product Name: SMAP (C11orf58) Human shRNA Lentiviral Particle (Locus ID 10944)

Locus ID:

chromosome 11 open reading frame 58; DKFZp686N0232; IMAGE145052; MGC117265; small Synonyms:

acidic protein; SMAP

Vector: pGFP-C-shLenti (TR30023)

Format: Lentiviral particles

C11orf58 - Human shRNA lentiviral particles (4 unique 29mer target-specific shRNA, 1 Components:

scramble control), 0.5 ml each, >10⁷ TU/ml.

RefSeq: NM 001142705, NM 014267, NM 014267.1, NM 014267.2, NM 014267.3, NM 014267.4,

NM 014267.5, NM 001142705.1, BC007103, BC016352, BC020937, BC103747, NM 014267.6

UniProt ID: 000193

These shRNA constructs were designed against multiple splice variants at this gene locus. To shRNA Design:

> be certain that your variant of interest is targeted, please contact <u>techsupport@origene.com</u>. If you need a special design or shRNA sequence, please utilize our custom shRNA service.

Performance

OriGene guarantees that the sequences in the shRNA expression cassettes are verified to **Guaranteed:** 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).

