## Product datasheet for TR504903

## Parn Mouse shRNA Plasmid (Locus ID 74108)

## Product data:

Product Type:
Product Name:
Locus ID:
Synonyms:
Vector:
E. coli Selection:

Mammalian Cell
Selection:
Format:
Components:

RefSeq:

UniProt ID:
Summary:

shRNA Plasmids

Parn Mouse shRNA Plasmid (Locus ID 74108)
74108
1200003I18Rik; DAN
pRS (TR20003)
Ampicillin
Puromycin

Retroviral plasmids
Parn - Mouse, 4 unique 29mer shRNA constructs in retroviral untagged vector(Gene ID = 74108). $5 \mu \mathrm{~g}$ purified plasmid DNA per construct

29-mer scrambled shRNA cassette in pRS Vector, TR30012, included for free.
BC021899, NM 028761, NM 001358452, NM 001358453, NM 028761.1 NM 028761.2, NM 028761.3

## Q8VDG3

3'-exoribonuclease that has a preference for $\operatorname{poly}(A)$ tails of mRNAs, thereby efficiently degrading poly $(A)$ tails. Exonucleolytic degradation of the poly $(A)$ tail is often the first step in the decay of eukaryotic mRNAs and is also used to silence certain maternal mRNAs translationally during oocyte maturation and early embryonic development. Interacts with both the 3'-end poly(A) tail and the 5'-end cap structure during degradation, the interaction with the cap structure being required for an efficient degradation of poly(A) tails. Involved in nonsense-mediated mRNA decay, a critical process of selective degradation of mRNAs that contain premature stop codons. Also involved in degradation of inherently unstable mRNAs that contain AU-rich elements (AREs) in their 3' UTR, possibly via its interaction with KHSRP. Probably mediates the removal of poly(A) tails of AREs mRNAs, which constitutes the first step of destabilization (By similarity). Also able to recognize poly(A) tails of microRNAs such as MIR21 and H/ACA box snoRNAs (small nucleolar RNAs) leading to leading to microRNAs degradation or snoRNA increased stability (By similarity).[UniProtKB/Swiss-Prot Function]
shRNA Design:

Performance Guaranteed:

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.

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).

