

Product datasheet for **TR503268**

Zbtb33 Mouse shRNA Plasmid (Locus ID 56805)

Product data:

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| Product Type: | shRNA Plasmids |
| Product Name: | Zbtb33 Mouse shRNA Plasmid (Locus ID 56805) |
| Locus ID: | 56805 |
| Synonyms: | AW260253; E130014G12Rik; Kaiso; Znf-kaiso |
| Vector: | pRS (TR20003) |
| E. coli Selection: | Ampicillin |
| Mammalian Cell Selection: | Puromycin |
| Format: | Retroviral plasmids |
| Components: | Zbtb33 - Mouse, 4 unique 29mer shRNA constructs in retroviral untagged vector(Gene ID = 56805). 5µg purified plasmid DNA per construct 29-mer scrambled shRNA cassette in pRS Vector, TR30012, included for free. |
| RefSeq: | NM_001079513 , NM_020256 , NM_020256.1 , NM_020256.2 , NM_001079513.1 , BC138647 , BC025136 , BC027282 , BC138648 |
| UniProt ID: | Q8BN78 |
| Summary: | Transcriptional regulator with bimodal DNA-binding specificity. Binds to methylated CpG dinucleotides in the consensus sequence 5'-CGCG-3' and also binds to the non-methylated consensus sequence 5'-CTGCNA-3' also known as the consensus kaiso binding site (KBS). May recruit the N-CoR repressor complex to promote histone deacetylation and the formation of repressive chromatin structures in target gene promoters. Contributes to the repression of target genes of the Wnt signaling pathway. May also activate transcription of a subset of target genes by the recruitment of CTNND2. Represses expression of MMP7 in conjunction with transcriptional corepressors CBFA2T3, CBFA2T2 and RUNX1T1 (By similarity). [UniProtKB/Swiss-Prot Function] |
| 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 . |



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**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).