

## Product datasheet for **TR503116**

### Acin1 Mouse shRNA Plasmid (Locus ID 56215)

#### Product data:

Product Type:	shRNA Plasmids
Product Name:	Acin1 Mouse shRNA Plasmid (Locus ID 56215)
Locus ID:	56215
Synonyms:	2610036I19Rik; 2610510L13Rik; Acinus; acinusL; acinusS; Acn; C79325; mKIAA0670
Vector:	pRS (TR20003)
E. coli Selection:	Ampicillin
Mammalian Cell Selection:	Puromycin
Format:	Retroviral plasmids
Components:	Acin1 - Mouse, 4 unique 29mer shRNA constructs in retroviral untagged vector(Gene ID = 56215). 5µg purified plasmid DNA per construct 29-mer scrambled shRNA cassette in pRS Vector, TR30012, included for free.
RefSeq:	<a href="#">NM_001085472</a> , <a href="#">NM_001085473</a> , <a href="#">NM_001242605</a> , <a href="#">NM_001242606</a> , <a href="#">NM_019567</a> , <a href="#">NM_023190</a> , <a href="#">NR_024029</a> , <a href="#">NR_024030</a> , <a href="#">NM_019567.1</a> , <a href="#">NM_019567.2</a> , <a href="#">NM_019567.3</a> , <a href="#">NM_001085473.1</a> , <a href="#">NM_001085473.2</a> , <a href="#">NM_023190.1</a> , <a href="#">NM_023190.2</a> , <a href="#">NM_023190.3</a> , <a href="#">NM_001085472.1</a> , <a href="#">NM_001085472.2</a> , <a href="#">NM_001242606.1</a> , <a href="#">NM_001242605.1</a> , <a href="#">BC019127</a> , <a href="#">BC052755</a> , <a href="#">BC075691</a> , <a href="#">BC094217</a> , <a href="#">BC169216</a> , <a href="#">BC169217</a>
UniProt ID:	<a href="#">Q9JIX8</a>



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<b>Summary:</b>	<p>Auxiliary component of the splicing-dependent multiprotein exon junction complex (EJC) deposited at splice junction on mRNAs. The EJC is a dynamic structure consisting of core proteins and several peripheral nuclear and cytoplasmic associated factors that join the complex only transiently either during EJC assembly or during subsequent mRNA metabolism. Component of the ASAP complexes which bind RNA in a sequence-independent manner and are proposed to be recruited to the EJC prior to or during the splicing process and to regulate specific excision of introns in specific transcription subsets; ACIN1 confers RNA-binding to the complex. The ASAP complex can inhibit RNA processing during in vitro splicing reactions. The ASAP complex promotes apoptosis and is disassembled after induction of apoptosis. Involved in the splicing modulation of BCL2L1/Bcl-X (and probably other apoptotic genes); specifically inhibits formation of proapoptotic isoforms such as Bcl-X(S); the activity is different from the established EJC assembly and function. Induces apoptotic chromatin condensation after activation by CASP3. Regulates cyclin A1, but not cyclin A2, expression in leukemia cells (By similarity).[UniProtKB/Swiss-Prot Function]</p>
<b>shRNA Design:</b>	<p>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 <a href="mailto:techsupport@origene.com">techsupport@origene.com</a>. If you need a special design or shRNA sequence, please utilize our <a href="#">custom shRNA service</a>.</p>
<b>Performance Guaranteed:</b>	<p>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.</p> <p>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 <a href="mailto:techsupport@origene.com">techsupport@origene.com</a>. Please provide your data indicating the transfection efficiency and measurement of gene expression knockdown compared to the scrambled shRNA control (Western Blot data preferred).</p>