

## Product datasheet for **TL313060**

### CD95 (FAS) Human shRNA Plasmid Kit (Locus ID 355)

#### Product data:

Product Type:	shRNA Plasmids
Product Name:	CD95 (FAS) Human shRNA Plasmid Kit (Locus ID 355)
Locus ID:	355
Synonyms:	ALPS1A; APO-1; APT1; CD95; FAS1; FASTM; TNFRSF6
Vector:	pGFP-C-shLenti (TR30023)
E. coli Selection:	Chloramphenicol (34 ug/ml)
Mammalian Cell Selection:	Puromycin
Format:	Lentiviral plasmids
Components:	FAS - Human, 4 unique 29mer shRNA constructs in lentiviral GFP vector(Gene ID = 355). 5µg purified plasmid DNA per construct 29-mer scrambled shRNA cassette in pGFP-C-shLenti Vector, TR30021, included for free.
RefSeq:	<a href="#">NM_000043</a> , <a href="#">NM_001320619</a> , <a href="#">NM_152871</a> , <a href="#">NM_152872</a> , <a href="#">NM_152873</a> , <a href="#">NM_152874</a> , <a href="#">NM_152875</a> , <a href="#">NM_152876</a> , <a href="#">NM_152877</a> , <a href="#">NR_028033</a> , <a href="#">NR_028034</a> , <a href="#">NR_028035</a> , <a href="#">NR_028036</a> , <a href="#">NR_135313</a> , <a href="#">NR_135314</a> , <a href="#">NR_135315</a> , <a href="#">NM_000043.1</a> , <a href="#">NM_000043.2</a> , <a href="#">NM_000043.3</a> , <a href="#">NM_000043.4</a> , <a href="#">NM_000043.5</a> , <a href="#">NM_152872.2</a> , <a href="#">NM_152872.3</a> , <a href="#">NM_152876.1</a> , <a href="#">NM_152871.1</a> , <a href="#">NM_152871.2</a> , <a href="#">NM_152871.3</a> , <a href="#">NM_152873.1</a> , <a href="#">NM_152875.1</a> , <a href="#">NM_152874.1</a> , <a href="#">NM_152877.1</a> , <a href="#">BC012479</a> , <a href="#">BC012479.1</a> , <a href="#">BC065736</a> , <a href="#">NM_152871.4</a> , <a href="#">NM_152872.4</a>
UniProt ID:	<a href="#">P25445</a>



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<b>Summary:</b>	<p>The protein encoded by this gene is a member of the TNF-receptor superfamily. This receptor contains a death domain. It has been shown to play a central role in the physiological regulation of programmed cell death, and has been implicated in the pathogenesis of various malignancies and diseases of the immune system. The interaction of this receptor with its ligand allows the formation of a death-inducing signaling complex that includes Fas-associated death domain protein (FADD), caspase 8, and caspase 10. The autoproteolytic processing of the caspases in the complex triggers a downstream caspase cascade, and leads to apoptosis. This receptor has been also shown to activate NF-kappaB, MAPK3/ERK1, and MAPK8/JNK, and is found to be involved in transducing the proliferating signals in normal diploid fibroblast and T cells. Several alternatively spliced transcript variants have been described, some of which are candidates for nonsense-mediated mRNA decay (NMD). The isoforms lacking the transmembrane domain may negatively regulate the apoptosis mediated by the full length isoform. [provided by RefSeq, Mar 2011]</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>