

## Product datasheet for **TR310508**

### PEG10 Human shRNA Plasmid Kit (Locus ID 23089)

#### Product data:

Product Type:	shRNA Plasmids
Product Name:	PEG10 Human shRNA Plasmid Kit (Locus ID 23089)
Locus ID:	23089
Synonyms:	EDR; HB-1; Mar2; Mart2; MEF3L; RGAG3; RTL2; SIRH1
Vector:	pRS (TR20003)
E. coli Selection:	Ampicillin
Mammalian Cell Selection:	Puromycin
Format:	Retroviral plasmids
Components:	PEG10 - Human, 4 unique 29mer shRNA constructs in retroviral untagged vector(Gene ID = 23089). 5µg purified plasmid DNA per construct 29-mer scrambled shRNA cassette in pRS Vector, TR30012, included for free.
RefSeq:	<a href="#">NM_001040152</a> , <a href="#">NM_001172437</a> , <a href="#">NM_001172438</a> , <a href="#">NM_001184961</a> , <a href="#">NM_001184962</a> , <a href="#">NM_015068</a> , <a href="#">NM_001040152.1</a> , <a href="#">NM_015068.1</a> , <a href="#">NM_015068.2</a> , <a href="#">NM_001184962.1</a> , <a href="#">NM_001172438.1</a> , <a href="#">NM_001172438.2</a> , <a href="#">NM_015068.3</a> , <a href="#">NM_001184961.1</a> , <a href="#">NM_001172437.1</a> , <a href="#">BC050659</a> , <a href="#">BC050659.2</a> , <a href="#">BC015448</a> , <a href="#">NM_001040152.2</a> , <a href="#">NM_001172438.3</a> , <a href="#">NM_001184962.2</a>
UniProt ID:	<a href="#">Q86TG7</a>



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<b>Summary:</b>	<p>This is a paternally expressed imprinted gene that is thought to have been derived from the Ty3/Gypsy family of retrotransposons. It contains two overlapping open reading frames, RF1 and RF2, and expresses two proteins: a shorter, gag-like protein (with a CCHC-type zinc finger domain) from RF1; and a longer, gag/pol-like fusion protein (with an additional aspartic protease motif) from RF1/RF2 by -1 translational frameshifting (-1 FS). While -1 FS has been observed in RNA viruses and transposons in both prokaryotes and eukaryotes, this gene represents the first example of -1 FS in a eukaryotic cellular gene. This gene is highly conserved across mammalian species and retains the heptanucleotide (GGGAAAC) and pseudoknot elements required for -1 FS. It is expressed in adult and embryonic tissues (most notably in placenta) and reported to have a role in cell proliferation, differentiation and apoptosis. Overexpression of this gene has been associated with several malignancies, such as hepatocellular carcinoma and B-cell lymphocytic leukemia. Knockout mice lacking this gene showed early embryonic lethality with placental defects, indicating the importance of this gene in embryonic development. Additional isoforms resulting from alternatively spliced transcript variants, and use of upstream non-AUG (CUG) start codon have been reported for this gene. [provided by RefSeq, Oct 2014]</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>