

## **Product datasheet for TR304007**

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## **HYAL1 Human shRNA Plasmid Kit (Locus ID 3373)**

**Product data:** 

**Product Type:** shRNA Plasmids

Product Name: HYAL1 Human shRNA Plasmid Kit (Locus ID 3373)

Locus ID: 3373

Synonyms: HYAL-1; LUCA1; MGC45987; NAT6

Vector: pRS (TR20003)

E. coli Selection: Ampicillin

Mammalian Cell Puromycin

Selection:

Format:

Retroviral plasmids

Components: HYAL1 - Human, 4 unique 29mer shRNA constructs in retroviral untagged vector(Gene ID =

3373). 5µg purified plasmid DNA per construct

29-mer scrambled shRNA cassette in pRS Vector, TR30012, included for free.

RefSeq: NM 007312, NM 033159, NM 153281, NM 153282, NM 153283, NM 153284, NM 153285,

NM 153286, NR 047690, NM 153285.1, NM 153285.2, NM 153282.1, NM 153282.2, NM 153281.1, NM 033159.1, NM 033159.2, NM 033159.3, NM 153283.1, NM 153283.2, NM 007312.3, NM 153284.1, NM 153286.1, BC035695, BC035695.1, BC025774, NM 033159.4,

NM 153282.3

UniProt ID: Q12794

Summary: This gene encodes a lysosomal hyaluronidase. Hyaluronidases intracellularly degrade

hyaluronan, one of the major glycosaminoglycans of the extracellular matrix. Hyaluronan is thought to be involved in cell proliferation, migration and differentiation. This enzyme is active at an acidic pH and is the major hyaluronidase in plasma. Mutations in this gene are associated with mucopolysaccharidosis type IX, or hyaluronidase deficiency. The gene is one

of several related genes in a region of chromosome 3p21.3 associated with tumor

suppression. Multiple transcript variants encoding different isoforms have been found for

this gene. [provided by RefSeq, Jul 2008]

**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 <u>techsupport@origene.com</u>. If you need a special design or shRNA sequence, please utilize our <u>custom shRNA service</u>.





## 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).