

## Product datasheet for **TL306992**

### ABHD5 Human shRNA Plasmid Kit (Locus ID 51099)

#### Product data:

Product Type:	shRNA Plasmids
Product Name:	ABHD5 Human shRNA Plasmid Kit (Locus ID 51099)
Locus ID:	51099
Synonyms:	CGI58; IECN2; NCIE2
Vector:	pGFP-C-shLenti (TR30023)
E. coli Selection:	Chloramphenicol (34 ug/ml)
Mammalian Cell Selection:	Puromycin
Format:	Lentiviral plasmids
Components:	ABHD5 - Human, 4 unique 29mer shRNA constructs in lentiviral GFP vector(Gene ID = 51099). 5µg purified plasmid DNA per construct 29-mer scrambled shRNA cassette in pGFP-C-shLenti Vector, TR30021, included for free.
RefSeq:	<a href="#">NM_016006</a> , <a href="#">NM_001355186</a> , <a href="#">NM_016006.1</a> , <a href="#">NM_016006.2</a> , <a href="#">NM_016006.3</a> , <a href="#">NM_016006.4</a> , <a href="#">BC021958</a> , <a href="#">NM_001365649</a> , <a href="#">NM_001365650</a> , <a href="#">NR_158560</a> , <a href="#">NM_016006.6</a>
UniProt ID:	<a href="#">Q8WTS1</a>
Summary:	The protein encoded by this gene belongs to a large family of proteins defined by an alpha/beta hydrolase fold, and contains three sequence motifs that correspond to a catalytic triad found in the esterase/lipase/thioesterase subfamily. It differs from other members of this subfamily in that its putative catalytic triad contains an asparagine instead of the serine residue. Mutations in this gene have been associated with Chanarin-Dorfman syndrome, a triglyceride storage disease with impaired long-chain fatty acid oxidation. [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 <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> .



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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](mailto: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).