

## OriGene Technologies, Inc.

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## Product datasheet for TL306506

## Protein atonal homolog 8 (ATOH8) Human shRNA Plasmid Kit (Locus ID 84913)

## **Product data:**

| Product Type:                | shRNA Plasmids   |
|------------------------------|--|
| Product Name:                | Protein atonal homolog 8 (ATOH8) Human shRNA Plasmid Kit (Locus ID 84913)  |
| Locus ID:                    | 84913  |
| Synonyms:                    | bHLHa21; HATH6   |
| Vector:                      | pGFP-C-shLenti (TR30023)   |
| E. coli Selection:           | Chloramphenicol (34 ug/ml)   |
| Mammalian Cell<br>Selection: | Puromycin  |
| Format:                      | Lentiviral plasmids  |
| Components:                  | ATOH8 - Human, 4 unique 29mer shRNA constructs in lentiviral GFP vector(Gene ID = 84913).<br>5μg purified plasmid DNA per construct<br>29-mer scrambled shRNA cassette in pGFP-C-shLenti Vector, TR30021, included for free. |
| RefSeq:                      | <u>NM 032827, NM 032827.1, NM 032827.2, NM 032827.3, NM 032827.4, NM 032827.5, BC019854, BC094832, BM671127</u>  |
| UniProt ID:                  | <u>Q96SQ7</u>  |



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| Transcription factor that binds a palindromic (canonical) core consensus DNA sequence 5'-<br>CANNTG- 3' known as an E-box element, possibly as a heterodimer with other bHLH proteins<br>(PubMed:24236640). Regulates endothelial cell proliferation, migration and tube-like<br>structures formation (PubMed:24463812). Modulates endothelial cell differentiation through<br>NOS3 (PubMed:24463812). May be implicated in specification and differentiation of neuronal<br>cell lineages in the brain (By similarity). May participate in kidney development and may be<br>involved in podocyte differentiation (By similarity). During early embryonic development is<br>involved in tissue-specific differentiation processes that are dependent on class II bHLH<br>factors and namely modulates the differentiation program initiated by the pro-endocrine<br>factor NEUROG3 (By similarity). During myogenesis, may play a role during the transition of<br>myoblasts from the proliferative phase to the differentiation phase (By similarity). Positively<br>regulates HAMP transcription in two ways, firstly by acting directly on the HAMP promoter via<br>E-boxes binding and indirectly through increased phosphorylation of SMAD protein complex<br>(PubMed:24236640). Repress NEUROG3-dependent gene activation in a gene-specific manner<br>through at least two mechanisms; requires only either the sequestering of a general partner<br>such as TCF3 through heterodimerization, either also requires binding of the bHLH domain to<br>DNA via a basic motif (By similarity).[UniProtKB/Swiss-Prot Function] |
| These shRNA constructs were designed against multiple splice variants at this gene locus. To<br>be certain that your variant of interest is targeted, please contact <u>techsupport@origene.com</u> .<br>If you need a special design or shRNA sequence, please utilize our <u>custom shRNA service</u> .  |
| 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.<br>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  |
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