

## Product datasheet for **TL308688**

### **p63 (TP63) Human shRNA Plasmid Kit (Locus ID 8626)**

#### **Product data:**

Product Type:	shRNA Plasmids
Product Name:	p63 (TP63) Human shRNA Plasmid Kit (Locus ID 8626)
Locus ID:	8626
Synonyms:	AIS; B(p51A); B(p51B); EEC3; KET; LMS; NBP; OFC8; p40; p51; p53CP; p63; p73H; p73L; RHS; SHFM4; TP53CP; TP53L; TP73L
Vector:	pGFP-C-shLenti (TR30023)
E. coli Selection:	Chloramphenicol (34 ug/ml)
Mammalian Cell Selection:	Puromycin
Format:	Lentiviral plasmids
Components:	TP63 - Human, 4 unique 29mer shRNA constructs in lentiviral GFP vector(Gene ID = 8626). 5µg purified plasmid DNA per construct 29-mer scrambled shRNA cassette in pGFP-C-shLenti Vector, TR30021, included for free.
RefSeq:	<a href="#">NM_001114978</a> , <a href="#">NM_001114979</a> , <a href="#">NM_001114980</a> , <a href="#">NM_001114981</a> , <a href="#">NM_001114982</a> , <a href="#">NM_001329144</a> , <a href="#">NM_001329145</a> , <a href="#">NM_001329146</a> , <a href="#">NM_001329148</a> , <a href="#">NM_001329149</a> , <a href="#">NM_001329150</a> , <a href="#">NM_001329964</a> , <a href="#">NM_003722</a> , <a href="#">NM_003722.1</a> , <a href="#">NM_003722.2</a> , <a href="#">NM_003722.3</a> , <a href="#">NM_003722.4</a> , <a href="#">NM_001114982.1</a> , <a href="#">NM_001114981.1</a> , <a href="#">NM_001114979.1</a> , <a href="#">NM_001114978.1</a> , <a href="#">NM_001114980.1</a> , <a href="#">BC039815</a> , <a href="#">BC039815.1</a> , <a href="#">NM_001114979.2</a> , <a href="#">NM_001114981.2</a> , <a href="#">NM_003722.5</a> , <a href="#">NM_001114980.2</a> , <a href="#">NM_001114978.2</a> , <a href="#">NM_001114982.2</a>
UniProt ID:	<a href="#">Q9H3D4</a>



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<b>Summary:</b>	<p>This gene encodes a member of the p53 family of transcription factors. The functional domains of p53 family proteins include an N-terminal transactivation domain, a central DNA-binding domain and an oligomerization domain. Alternative splicing of this gene and the use of alternative promoters results in multiple transcript variants encoding different isoforms that vary in their functional properties. These isoforms function during skin development and maintenance, adult stem/progenitor cell regulation, heart development and premature aging. Some isoforms have been found to protect the germline by eliminating oocytes or testicular germ cells that have suffered DNA damage. Mutations in this gene are associated with ectodermal dysplasia, and cleft lip/palate syndrome 3 (EEC3); split-hand/foot malformation 4 (SHFM4); ankyloblepharon-ectodermal defects-cleft lip/palate; ADULT syndrome (acrodermato-ungual-lacrimal-tooth); limb-mammary syndrome; Rap-Hodgkin syndrome (RHS); and orofacial cleft 8. [provided by RefSeq, Aug 2016]</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>