

## Product datasheet for **TR314562**

### ATP7A Human shRNA Plasmid Kit (Locus ID 538)

#### Product data:

Product Type:	shRNA Plasmids
Product Name:	ATP7A Human shRNA Plasmid Kit (Locus ID 538)
Locus ID:	538
Synonyms:	DSMAX; MK; MNK; SMAX3
Vector:	pRS (TR20003)
E. coli Selection:	Ampicillin
Mammalian Cell Selection:	Puromycin
Format:	Retroviral plasmids
Components:	ATP7A - Human, 4 unique 29mer shRNA constructs in retroviral untagged vector(Gene ID = 538). 5µg purified plasmid DNA per construct 29-mer scrambled shRNA cassette in pRS Vector, TR30012, included for free.
RefSeq:	<a href="#">NM_000052</a> , <a href="#">NM_001282224</a> , <a href="#">NR_104109</a> , <a href="#">NM_000052.1</a> , <a href="#">NM_000052.2</a> , <a href="#">NM_000052.3</a> , <a href="#">NM_000052.4</a> , <a href="#">NM_000052.5</a> , <a href="#">NM_000052.6</a> , <a href="#">NM_001282224.1</a> , <a href="#">BC156437</a>
UniProt ID:	<a href="#">Q04656</a>
Summary:	This gene encodes a transmembrane protein that functions in copper transport across membranes. This protein is localized to the trans Golgi network, where it is predicted to supply copper to copper-dependent enzymes in the secretory pathway. It relocalizes to the plasma membrane under conditions of elevated extracellular copper, and functions in the efflux of copper from cells. Mutations in this gene are associated with Menkes disease, X-linked distal spinal muscular atrophy, and occipital horn syndrome. Alternatively-spliced transcript variants have been observed. [provided by RefSeq, Aug 2013]
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).