

## Product datasheet for **TL314261**

### Carbonic Anhydrase I (CA1) Human shRNA Plasmid Kit (Locus ID 759)

#### Product data:

Product Type:	shRNA Plasmids
Product Name:	Carbonic Anhydrase I (CA1) Human shRNA Plasmid Kit (Locus ID 759)
Locus ID:	759
Synonyms:	CA-I; CAB; Car1; HEL-S-11
Vector:	pGFP-C-shLenti (TR30023)
E. coli Selection:	Chloramphenicol (34 ug/ml)
Mammalian Cell Selection:	Puromycin
Format:	Lentiviral plasmids
Components:	CA1 - Human, 4 unique 29mer shRNA constructs in lentiviral GFP vector(Gene ID = 759). 5µg purified plasmid DNA per construct 29-mer scrambled shRNA cassette in pGFP-C-shLenti Vector, TR30021, included for free.
RefSeq:	<a href="#">NM_001128829</a> , <a href="#">NM_001128830</a> , <a href="#">NM_001128831</a> , <a href="#">NM_001164830</a> , <a href="#">NM_001291967</a> , <a href="#">NM_001291968</a> , <a href="#">NM_001738</a> , <a href="#">NM_001738.1</a> , <a href="#">NM_001738.2</a> , <a href="#">NM_001738.3</a> , <a href="#">NM_001738.4</a> , <a href="#">NM_001128829.1</a> , <a href="#">NM_001128829.2</a> , <a href="#">NM_001128829.3</a> , <a href="#">NM_001128830.1</a> , <a href="#">NM_001128830.2</a> , <a href="#">NM_001128830.3</a> , <a href="#">NM_001128831.1</a> , <a href="#">NM_001128831.2</a> , <a href="#">NM_001128831.3</a> , <a href="#">NM_001164830.1</a> , <a href="#">NM_001291968.1</a> , <a href="#">NM_001291967.1</a> , <a href="#">BC027890</a> , <a href="#">BC027890.1</a> , <a href="#">NM_001738.5</a> , <a href="#">NM_001291968.2</a> , <a href="#">NM_001128829.4</a> , <a href="#">NM_001164830.2</a> , <a href="#">NM_001128830.4</a> , <a href="#">NM_001291967.2</a> , <a href="#">NM_001128831.4</a>
UniProt ID:	<a href="#">P00915</a>
Summary:	Carbonic anhydrases (CAs) are a large family of zinc metalloenzymes that catalyze the reversible hydration of carbon dioxide. They participate in a variety of biological processes, including respiration, calcification, acid-base balance, bone resorption, and the formation of aqueous humor, cerebrospinal fluid, saliva and gastric acid. They show extensive diversity in tissue distribution and in their subcellular localization. This CA1 gene is closely linked to the CA2 and CA3 genes on chromosome 8. It encodes a cytosolic protein that is found at the highest level in erythrocytes. Allelic variants of this gene have been described in some populations. Alternative splicing and the use of alternative promoters results in multiple transcript variants. [provided by RefSeq, Nov 2016]



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**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 [techsupport@origene.com](mailto:techsupport@origene.com). If you need a special design or shRNA sequence, please utilize our [custom shRNA service](#).

**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).