

## Product datasheet for **TL314051**

### CDC27 Human shRNA Plasmid Kit (Locus ID 996)

#### Product data:

|                           |  |
|---------------------------|--|
| Product Type:             | shRNA Plasmids   |
| Product Name:             | CDC27 Human shRNA Plasmid Kit (Locus ID 996)   |
| Locus ID:                 | 996  |
| Synonyms:                 | ANAPC3; APC3; CDC27Hs; D0S1430E; D17S978E; H-NUC; HNUC; NUC2   |
| Vector:                   | pGFP-C-shLenti (TR30023)   |
| E. coli Selection:        | Chloramphenicol (34 ug/ml)   |
| Mammalian Cell Selection: | Puromycin  |
| Format:                   | Lentiviral plasmids  |
| Components:               | CDC27 - Human, 4 unique 29mer shRNA constructs in lentiviral GFP vector(Gene ID = 996).<br>5µg purified plasmid DNA per construct<br>29-mer scrambled shRNA cassette in pGFP-C-shLenti Vector, TR30021, included for free.   |
| RefSeq:                   | <a href="#">NM_001114091</a> , <a href="#">NM_001256</a> , <a href="#">NM_001293089</a> , <a href="#">NM_001293091</a> , <a href="#">NM_001353035</a> ,<br><a href="#">NM_001353047</a> , <a href="#">NM_001353049</a> , <a href="#">NM_001353050</a> , <a href="#">NM_001353051</a> , <a href="#">NR_148340</a> , <a href="#">NM_001256.1</a> ,<br><a href="#">NM_001256.2</a> , <a href="#">NM_001256.3</a> , <a href="#">NM_001114091.1</a> , <a href="#">NM_001114091.2</a> , <a href="#">NM_001293091.1</a> ,<br><a href="#">NM_001293089.1</a> , <a href="#">BC011656</a> , <a href="#">NM_001293091.3</a> , <a href="#">NM_001114091.4</a> , <a href="#">NM_001293089.3</a> ,<br><a href="#">NM_001256.6</a>  |
| UniProt ID:               | <a href="#">P30260</a>   |
| Summary:                  | The protein encoded by this gene shares strong similarity with <i>Saccharomyces cerevisiae</i> protein Cdc27, and the gene product of <i>Schizosaccharomyces pombe</i> nuc 2. This protein is a component of the anaphase-promoting complex (APC), which is composed of eight protein subunits and is highly conserved in eukaryotic cells. This complex catalyzes the formation of cyclin B-ubiquitin conjugate, which is responsible for the ubiquitin-mediated proteolysis of B-type cyclins. The protein encoded by this gene and three other members of the APC complex contain tetratricopeptide (TPR) repeats, which are important for protein-protein interactions. This protein was shown to interact with mitotic checkpoint proteins including Mad2, p55CDC and BUBR1, and it may thus be involved in controlling the timing of mitosis. Alternative splicing of this gene results in multiple transcript variants. Related pseudogenes have been identified on chromosomes 2, 22 and Y. [provided by RefSeq, May 2014] |



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|--------------------------------|---|
| <b>shRNA Design:</b>           | 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> .  |
| <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> |