

## Product datasheet for **TL314008**

### CEACAM1 Human shRNA Plasmid Kit (Locus ID 634)

#### Product data:

|                           |   |
|---------------------------|---|
| Product Type:             | shRNA Plasmids  |
| Product Name:             | CEACAM1 Human shRNA Plasmid Kit (Locus ID 634)  |
| Locus ID:                 | 634   |
| Synonyms:                 | BGP; BGP1; BGPI   |
| Vector:                   | pGFP-C-shLenti (TR30023)  |
| E. coli Selection:        | Chloramphenicol (34 ug/ml)  |
| Mammalian Cell Selection: | Puromycin   |
| Format:                   | Lentiviral plasmids   |
| Components:               | CEACAM1 - Human, 4 unique 29mer shRNA constructs in lentiviral GFP vector(Gene ID = 634).<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_001024912</a> , <a href="#">NM_001184813</a> , <a href="#">NM_001184815</a> , <a href="#">NM_001184816</a> , <a href="#">NM_001205344</a> ,<br><a href="#">NM_001712</a> , <a href="#">NM_001024912.1</a> , <a href="#">NM_001024912.2</a> , <a href="#">NM_001712.1</a> , <a href="#">NM_001712.2</a> , <a href="#">NM_001712.3</a> ,<br><a href="#">NM_001712.4</a> , <a href="#">NM_001184816.1</a> , <a href="#">NM_001184813.1</a> , <a href="#">NM_001184815.1</a> , <a href="#">NM_001205344.1</a> ,<br><a href="#">BC014473</a> , <a href="#">BC014473.1</a> , <a href="#">BC024164</a> , <a href="#">NM_001184813.2</a> , <a href="#">NM_001205344.2</a> , <a href="#">NM_001184815.2</a> ,<br><a href="#">NM_001712.5</a> , <a href="#">NM_001184816.2</a> |
| UniProt ID:               | <a href="#">P13688</a>  |



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**Summary:**

This gene encodes a member of the carcinoembryonic antigen (CEA) gene family, which belongs to the immunoglobulin superfamily. Two subgroups of the CEA family, the CEA cell adhesion molecules and the pregnancy-specific glycoproteins, are located within a 1.2 Mb cluster on the long arm of chromosome 19. Eleven pseudogenes of the CEA cell adhesion molecule subgroup are also found in the cluster. The encoded protein was originally described in bile ducts of liver as biliary glycoprotein. Subsequently, it was found to be a cell-cell adhesion molecule detected on leukocytes, epithelia, and endothelia. The encoded protein mediates cell adhesion via homophilic as well as heterophilic binding to other proteins of the subgroup. Multiple cellular activities have been attributed to the encoded protein, including roles in the differentiation and arrangement of tissue three-dimensional structure, angiogenesis, apoptosis, tumor suppression, metastasis, and the modulation of innate and adaptive immune responses. Multiple transcript variants encoding different isoforms have been reported, but the full-length nature of all variants has not been defined. [provided by RefSeq, May 2010]

**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).