

Product datasheet for **TR306361**

BSCL2 Human shRNA Plasmid Kit (Locus ID 26580)

Product data:

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| Product Type: | shRNA Plasmids |
| Product Name: | BSCL2 Human shRNA Plasmid Kit (Locus ID 26580) |
| Locus ID: | 26580 |
| Synonyms: | GNG3LG; HMN5; HMN5C; PELD; SPG17 |
| Vector: | pRS (TR20003) |
| E. coli Selection: | Ampicillin |
| Mammalian Cell Selection: | Puromycin |
| Format: | Retroviral plasmids |
| Components: | BSCL2 - Human, 4 unique 29mer shRNA constructs in retroviral untagged vector(Gene ID = 26580). 5µg purified plasmid DNA per construct 29-mer scrambled shRNA cassette in pRS Vector, TR30012, included for free. |
| RefSeq: | NM_001122955 , NM_001130702 , NM_032667 , NR_037948 , NR_037949 , NM_032667.2 , NM_032667.3 , NM_032667.4 , NM_032667.5 , NM_032667.6 , NM_001130702.1 , NM_001130702.2 , NM_001122955.1 , NM_001122955.2 , NM_001122955.3 , BC009866 , BC093048 , BC004911 , BC012140 , BC041640 , BM763152 |
| UniProt ID: | Q96G97 |
| Summary: | This gene encodes the multi-pass transmembrane protein protein seipin. This protein localizes to the endoplasmic reticulum and may be important for lipid droplet morphology. Mutations in this gene have been associated with congenital generalized lipodystrophy type 2 or Berardinelli-Seip syndrome, a rare autosomal recessive disease characterized by a near absence of adipose tissue and severe insulin resistance. Alternatively spliced transcript variants encoding different isoforms have been found for this gene. Naturally occurring read-through transcription occurs between this locus and the neighboring locus HNRNPUL2 (heterogeneous nuclear ribonucleoprotein U-like 2).[provided by RefSeq, Mar 2011] |
| 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 . If you need a special design or shRNA sequence, please utilize our custom shRNA service . |



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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. Please provide your data indicating the transfection efficiency and measurement of gene expression knockdown compared to the scrambled shRNA control (Western Blot data preferred).