

Product datasheet for **TL313645**

CUG BP1 (CELF1) Human shRNA Plasmid Kit (Locus ID 10658)

Product data:

Product Type:	shRNA Plasmids
Product Name:	CUG BP1 (CELF1) Human shRNA Plasmid Kit (Locus ID 10658)
Locus ID:	10658
Synonyms:	BRUNOL2; CUG-BP; CUGBP; CUGBP1; EDEN-BP; hNab50; NAB50; NAPOR
Vector:	pGFP-C-shLenti (TR30023)
E. coli Selection:	Chloramphenicol (34 ug/ml)
Mammalian Cell Selection:	Puromycin
Format:	Lentiviral plasmids
Components:	CELF1 - Human, 4 unique 29mer shRNA constructs in lentiviral GFP vector(Gene ID = 10658). 5µg purified plasmid DNA per construct 29-mer scrambled shRNA cassette in pGFP-C-shLenti Vector, TR30021, included for free.
RefSeq:	NM_001025596 , NM_001172639 , NM_001172640 , NM_006560 , NM_198700 , NM_001330272 , NM_006560.1 , NM_006560.2 , NM_006560.3 , NM_198700.1 , NM_198700.2 , NM_001025596.1 , NM_001025596.2 , NM_001172640.1 , NM_001172639.1 , BC031079 , BC031079.1 , BC036782 , BM737883 , NM_001172640.2 , NM_198700.3 , NM_001172639.2 , NM_001025596.3 , NM_006560.4
UniProt ID:	Q92879
Summary:	Members of the CELF/BRUNOL protein family contain two N-terminal RNA recognition motif (RRM) domains, one C-terminal RRM domain, and a divergent segment of 160-230 aa between the second and third RRM domains. Members of this protein family regulate pre-mRNA alternative splicing and may also be involved in mRNA editing, and translation. This gene may play a role in myotonic dystrophy type 1 (DM1) via interactions with the dystrophia myotonica-protein kinase (DMPK) gene. Alternative splicing results in multiple transcript variants encoding different isoforms. [provided by RefSeq, Jul 2008]
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).