

Product datasheet for **RC206275L1V**

CUG BP1 (CELF1) (NM_006560) Human Tagged ORF Clone Lentiviral Particle

Product data:

| | |
|---------------------------|--|
| Product Type: | Lentiviral Particles |
| Product Name: | CUG BP1 (CELF1) (NM_006560) Human Tagged ORF Clone Lentiviral Particle |
| Symbol: | CUG BP1 |
| Synonyms: | BRUNOL2; CUG-BP; CUGBP; CUGBP1; EDEN-BP; hNab50; NAB50; NAPOR |
| Mammalian Cell Selection: | None |
| Vector: | pLenti-C-Myc-DDK (PS100064) |
| Tag: | Myc-DDK |
| ACCN: | NM_006560 |
| ORF Size: | 1449 bp |
| ORF Nucleotide Sequence: | The ORF insert of this clone is exactly the same as(RC206275). |
| OTI Disclaimer: | The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing variants is recommended prior to use. More info |
| OTI Annotation: | This clone was engineered to express the complete ORF with an expression tag. Expression varies depending on the nature of the gene. |
| RefSeq: | NM_006560.2 |
| RefSeq Size: | 4711 bp |
| RefSeq ORF: | 1449 bp |
| Locus ID: | 10658 |
| UniProt ID: | Q92879 |
| Cytogenetics: | 11p11.2 |
| Domains: | RRM |
| Protein Families: | Druggable Genome |



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MW: 51.6 kDa

Gene 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 dystrophin myotonia-protein kinase (DMPK) gene. Alternative splicing results in multiple transcript variants encoding different isoforms. [provided by RefSeq, Jul 2008]