

## Product datasheet for **MC216627**

### Celf2 (NM\_001110232) Mouse Untagged Clone

#### Product data:

|                           |   |
|---------------------------|---|
| Product Type:             | Expression Plasmids   |
| Product Name:             | Celf2 (NM_001110232) Mouse Untagged Clone   |
| Tag:                      | Tag Free  |
| Symbol:                   | Celf2   |
| Synonyms:                 | B230218O03; B230345P09Rik; C88023; CELF-2; CUG-BP2; Cugbp2; D230046B21Rik; Etr-3; mETR-3; Napor |
| Mammalian Cell Selection: | Neomycin  |
| Vector:                   | pCMV6-Entry (PS100001)  |
| E. coli Selection:        | Kanamycin (25 ug/mL)  |



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**Fully Sequenced ORF:** >MC216627 representing NM\_001110232  
 Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC  
 GCC**CGGATCGCC**

ATGAACGGAGCTTTGGATCATTAGACAGCCAGACCCAGATGCCATTAAGATGTTTGTGCGACAGATCC  
 CTAGGTCCTGGTCGAAAAGGAGCTGAAAGAACTTTTGGAGCCTTATGGAGCTGTCTACCAGATCAACGT  
 CCTCCGGGACCGAGTCAGAACCTCCCAAGAGTAAAGGTTGTTGTTTCGTAACATTTTATACAAGAAAA  
 GCTGCACTTGAGGCCAGAAATGCACTGCACAATATTAACCTTTACCTGGGATGCATCATCCATTCCAGA  
 TGAAACCTGCAGATAGTGAAGTCAAACGCTGTGGAAGACAGAAAATTGTTTCATAGGAATGGTTTCCAA  
 GAAATGTAAACGAGAATGATATCAGAGTGATGTTTTCTCCATTCGGTCAGATAGAAGAAATGCCGATTCTC  
 CGGGGACCTGATGGGCTGAGTCGAGGCTGTGCGTTTGTACATTTTCTACAAGGGCAATGCCACAGAAATG  
 CAATCAAAGCCATGCATCAGTCTCAGACCATGGAGGGTGTCTTCCACCAATCGTGGTGAAGTTTGCTGA  
 CACTCAGAAGGACAAAGAGCAAAGGCGCCTCCAGCAGCAGCTTGACAGCAGATGCAACAGCTCAACACT  
 GCCACTTGGGGGAACCTAACAGGACTGGGTGGACTTACCCCGCAGTACCTGGCGCTTCTGCAGCAGGCCA  
 CCTCTCCAGCAACCTGGGTGCATTAGTGGCATTAGCAAAATGGCTGGCATGAATGCTTTACAGTTACA  
 GAATCTGGCAACACTGGCTGCTGCTGCAGCTGCTGCTCAAACCTCAGCCACCAGCAACATGCAACCCCT  
 CTGTCTAGCACAAGCAGTGCCCTGGGAGCCCTCACAAGCCCTGTGGCTGCTTCAACCCCCAATTCACCG  
 CTGGTGCGGCCATGAATTCCTTGACCTCTCTGGGACTCTACAAGGATTGGCTGGAGCCACTGTCGGATT  
 GAATAATATTAATGCACTAGCAGGTATGGCGGCTCTGAATGGAGGACTTGGCGCCACAGGCTTGACGAAT  
 GGTACGGCTGGCACCATGGACGCCCTGACCCAGGCCTACTCAGGAATTCAGCAGTATGCGGCAGCTGCAC  
 TGCCCACTTTGTACGCCAGAGCTTGCTGCAACAGCAGAGTGCTGCAGGCAGCCAGAAGGAAGTCCAGA  
 GGGGGCAAACCTCTTTATTTACCACCTCCACAGGAGTTTGGAGACCAGGACATTCTGCAGATGTTTCATG  
 CCCTTTGGAAATGTTATCTCTGCTAAAGTCTTCATTGACAAACAGACCAATCTGAGCAAGTGCTTTGGTT  
 TTGTTAGCTATGACAATCCAGTCTCTGCACAAGCTGCAATCCAGGCTATGAACGGCTTTAGATCGGCAT  
 GAAACGCTTGAAGGTGCAGCTGAAACGCTCCAAAAACGACAGCAAACCTTACT**GA**

**ACGCGT**ACGCGGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT  
 ACAAGGATGACGACGATAAGGTTTAA

**Restriction Sites:** Sgfl-MluI

**ACCN:** NM\_001110232

**Insert Size:** 1455 bp

**OTI Disclaimer:** Our molecular clone sequence data has been matched to the reference identifier above as a point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative RNA splicing form or single nucleotide polymorphism (SNP).

**Components:** The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).

**Reconstitution Method:**

1. Centrifuge at 5,000xg for 5min.
2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.
3. Close the tube and incubate for 10 minutes at room temperature.
4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.
5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of shipping when stored at -20°C.

|                      |   |
|----------------------|---|
| <b>Note:</b>         | Plasmids are not sterile. For experiments where strict sterility is required, filtration with 0.22um filter is required.  |
| <b>RefSeq:</b>       | <a href="#">NM_001110232.1</a> , <a href="#">NP_001103702.1</a>   |
| <b>RefSeq Size:</b>  | 7735 bp   |
| <b>RefSeq ORF:</b>   | 1455 bp   |
| <b>Locus ID:</b>     | 14007   |
| <b>UniProt ID:</b>   | <a href="#">Q9Z0H4</a>  |
| <b>Cytogenetics:</b> | 2 A1  |
| <b>Gene Summary:</b> | <p>RNA-binding protein implicated in the regulation of several post-transcriptional events. Involved in pre-mRNA alternative splicing, mRNA translation and stability. Mediates exon inclusion and/or exclusion in pre-mRNA that are subject to tissue-specific and developmentally regulated alternative splicing (By similarity). Specifically activates exon 5 inclusion of TNNT2 in embryonic, but not adult, skeletal muscle (By similarity). Activates TNNT2 exon 5 inclusion by antagonizing the repressive effect of PTB (By similarity). Acts as both an activator and repressor of a pair of coregulated exons: promotes inclusion of the smooth muscle (SM) exon but exclusion of the non-muscle (NM) exon in actinin pre-mRNAs (By similarity). Promotes inclusion of exonS 21 and exclusion of exon 5 of the NMDA receptor R1 pre-mRNA (By similarity). Involved in the apoB RNA editing activity (By similarity). Increases COX2 mRNA stability and inhibits COX2 mRNA translation in epithelial cells after radiation injury. Modulates the cellular apoptosis program by regulating COX2-mediated prostaglandin E2 (PGE2) expression. Binds to (CUG)<sub>n</sub> triplet repeats in the 3' UTR of transcripts such as DMPK (By similarity). Binds to the muscle-specific splicing enhancer (MSE) intronic sites flanking the TNNT2 alternative exon 5 (By similarity). Binds preferentially to UG-rich sequences, in particular UG repeat and UGUU motifs (By similarity). Binds to apoB mRNA, specifically to AU-rich sequences located immediately upstream of the edited cytidine (By similarity). Binds AU-rich sequences in the 3' UTR of COX2 mRNA. Binds to an intronic RNA element responsible for the silencing of exon 21 splicing. Binds to (CUG)<sub>n</sub> repeats. May be a specific regulator of miRNA biogenesis. Binds to primary microRNA pri-MIR140 and, with CELF1, negatively regulates the processing to mature miRNA (By similarity).[UniProtKB/Swiss-Prot Function]</p> <p>Transcript Variant: This variant (5) differs in the 5' UTR and has multiple differences in the presence and absence of exons at its 5' end, compared to variant 1. These differences cause translation initiation at a downstream AUG and an isoform (5) that is shorter than isoform 1.</p> <p>Sequence Note: This RefSeq record was created from transcript and genomic sequence data to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on transcript alignments.</p> |