

## Product datasheet for MC208328

### Crem (NM\_001110859) Mouse Untagged Clone

#### Product data:

**Product Type:** Expression Plasmids  
**Product Name:** Crem (NM\_001110859) Mouse Untagged Clone  
**Tag:** Tag Free  
**Symbol:** Crem  
**Synonyms:** IC; ICER; ICERI  
**Vector:** pCMV6-Entry (PS100001)  
**E. coli Selection:** Kanamycin (25 ug/mL)  
**Cell Selection:** Neomycin  
**Fully Sequenced ORF:** >MC208328 representing NM\_001110859  
 Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC  
 GCCGCGATCGCC

ATGAGCAAATGTGGCAGGAAAAAGTATATGAGGACAAATGTAAGGCAAATGACCATGGAACAGTTGAAT  
 CACAGCAGGATCGAAGTGAACACGTTCTGTGGCAGAGCATAGCTCTGCTCATATGCAGACTGGTCAAAT  
 TTCTGTTCTACTCTAGCTCAGTTTCTGTAGCTGGATCAGGCACTGGAAGAGGCTCCCCAGCTGTGACT  
 CTAGTACAGTTACCTTCAGGCCAACTGTACAGGTCCAGGGAGTTATTCAGACACCACATCCATCGGTTA  
 TTCAATCACCAAAATACAACTGTTTCAGGTAGCAAAATTGCAGAGACAGATGATTCTGCAGACTCAGA  
 AGTAATTGATTCCGATAAACGTAGAGAAATTTTTCACGAAGACCCTCATATAGAAAAATACTGAATGAA  
 CTTTCTCTGTGTGCCTGGTATCCCAAGATTGAAGAAGAAAAATCAGAGGAAGAAGGGACACCACCTA  
 ACATTGCTACCATGGCAGTACCAACTAGCATATATCAGACTAGCACGGGGCAATACATTGCTATAGCTCA  
 AGGTGGAACAATCCAGATTTCTAACCCAGGATCTGATGGTGTTCAGGGACTCCAGGCATTAACAATGACA  
 AATTCAGGAGCTCCTCCGCCAGGTGCTACAATTGTACAGTATGCAGCACAATCAGCCGATGGTACACAGC  
 AGTTCTTTGTCCCAGGCAGCCAGTTGTTGTTCAAGATGAGGAGACTGACCTTGCCCAAGGTGTGGTG  
 TGCTGCCACAGGTGACATGCCAACTTACCAGATCCGAGCTCCTACTACTGCTTTGCCACAAGGTGTGGTG  
 ATGGCTGCCTCACAGGAAGCCTGCACAGTCCCCAGCAACTAGCAGAAAGCAACTCGCAAGCGGGAGC  
 TGAGGCTGATGAAAAACAGGGAAGCTGCTAAAGAATGTCGACGTCGAAAGAAAGAGTATGTGAAGTGTCT  
 TGAGAGTCGAGTCGAGTCTGGAAGTTGAGAACAAGAAGCTTATAGAGGAGCTTGAACTTTGAAAGAC  
 ATTTGCTCTCCCAAAACAGATTAG

ACGCGTACGCGGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT  
 ACAAGGATGACGACGATAAGGTTTAA

**Restriction Sites:** SgfI-MluI  
**ACCN:** NM\_001110859



<b>Insert Size:</b>	1074 bp
<b>OTI Disclaimer:</b>	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).
<b>Components:</b>	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).
<b>Reconstitution Method:</b>	<ol style="list-style-type: none"><li>1. Centrifuge at 5,000xg for 5min.</li><li>2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.</li><li>3. Close the tube and incubate for 10 minutes at room temperature.</li><li>4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.</li><li>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.</li></ol>
<b>RefSeq:</b>	<u><a href="#">NM_001110859.2</a></u> , <u><a href="#">NP_001104329.1</a></u>
<b>RefSeq Size:</b>	2362 bp
<b>RefSeq ORF:</b>	1074 bp
<b>Locus ID:</b>	12916
<b>UniProt ID:</b>	<u><a href="#">P27699</a></u>
<b>Cytogenetics:</b>	18 A1
<b>Gene Summary:</b>	<p>This gene encodes a basic-leucine zipper domain-containing protein that localizes to gene promoters, where it binds to the cyclic AMP response element (CRE). Different protein isoforms encoded by this gene may function as either activators or repressors of transcription. Activity of this gene is important in multiple developmental processes, including spermatogenesis. Mutation of this gene causes male infertility. Alternative splicing and promoter usage result in multiple transcript variants for this gene. [provided by RefSeq, Oct 2012]</p> <p>Transcript Variant: This variant (1, also known as tau) encodes the longest isoform (1). The structure of this variant is described in PMID: 1370576. Sequence Note: The RefSeq transcript and protein were derived from genomic sequence to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on alignments.</p>