

Product datasheet for **MC229708**

Cacna1c (NM_001256001) Mouse Untagged Clone

Product data:

Product Type: Expression Plasmids
Product Name: Cacna1c (NM_001256001) Mouse Untagged Clone
Tag: Tag Free
Symbol: Cacna1c
Synonyms: Cav1.2; Cchl1a1; D930026N18Rik; MBC; MELC-CC
Vector: pCMV6-Entry (PS100001)
E. coli Selection: Kanamycin (25 ug/mL)
Cell Selection: Neomycin
Fully Sequenced ORF: >MC229708 representing NM_001256001
Red=Cloning site **Blue**=ORF **Orange**=Stop codon

TTTTGTAATACGACTCACTATAGGGCGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
GCC**CGGATCGCC**

ATGGTCAATGAAAACACGAGGATGTACGTTCCAGAGGAAAACCACCAAGGTTCCAACATATGGGAGCCAC
GCCAGCTCATGCCAACATGAATGCCAATGCAGCTGCAGGACTTGCTCCCAGCACATCCCTACTCCAGG
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GCACAACCGCCACACGGCCGCCCGGGCTGCTGTGTCTGACCCTGAAGAACCCTATCCGGAGGGCGTG
CATAAGCATTGTTGAATGGAAACATTTGAAATCATTATTTTACTGACTATTTTGGCAATTGTGTGGCC
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 TGAAGCTGCTGAGCCGCGGGGAAGGCATCCGAACCCTGCTGTGGACCTTCAAGTCTTCCAGGCTCT
 GCCCTATGTGGCTCTTTGATTGTGATGCTGTTCTTTATCTATGCAGTATTGGGATCAGGTGTTTGGG
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 GCTACTCATCCACGGGCTCCAATGCCAACATCAACAATGCCAACAACTGCCCTGGGCCGCTTCCCCCA
 TCCCGTGGTACTCCAGCACGGTCAGCACTGTGGAGGGCCATGGGCCTCCCTTGTCCCTGTGTCCGA
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 CTTGCATCTGGTTCATCATCAGGCATTGGCAGTGGCAGGCTTGAGCCCCCTCCTGCAGAGAAGCCATTCT
 CCTACCACATTCCCAGGCCGTGCCACACCCCTGTCACTCCAGGCAGCCGGGCAGACCCCTACGGC
 CCATCCCTACCCTACGGCTGGAGGGGCGAGTCCAGCGAGAACTCAACAGCAGCTTCCATCCATCCA
 CTGCAGCTCCTGGTCTGAGGAGACGACAGCCTGTAGTGGGAGCAGCAGCATGGCCCGAGAGCCCGGCC
 GTCTCCCTCACCGTGCCAGCCAGGCTGGAGCTCCAGGGAGACAGTTCCATGGCAGTGCCAGCAGCTGG
 TGAAGCGGTCTTGATTTAGAAGGACTGGGACAGTTTGTCAAGATCCCAAGTTCATCGAGGTCACCAC
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 GGGGGCGCCAGCAGAGCCCAACCGCACCCCTTACCTTTTGTGAAGTGCAGGGACCCGGGCAGGACA
 GGGCTGTGGCCCCAGAGGACGAGAGCTGCGCATATGCCCTGGGGCGAGCCGGAGCGAGGAGCGCTCGC
 GGACAGCAGGTCTACGTCAGCAACCTGTAG

ACGCGTACGCGGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT
 ACAAGGATGACGACGATAAGGTTTAA

- Restriction Sites:** SgfI-MluI
- ACCN:** NM_001256001
- Insert Size:** 6471 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).
- OTI Annotation:** Clone contains native stop codon, and expresses the complete ORF without any c-terminal tag.
- 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.
- RefSeq:** [NM_001256001.2](#), [NP_001242930.1](#)
- RefSeq Size:** 13643 bp
- RefSeq ORF:** 6471 bp

Locus ID: 12288

Cytogenetics: 6 55.86 cM

Gene Summary: Pore-forming, alpha-1C subunit of the voltage-gated calcium channel that gives rise to L-type calcium currents (PubMed:14609949, PubMed:18586882, PubMed:21216955, PubMed:25368181, PubMed:28119464). Mediates influx of calcium ions into the cytoplasm, and thereby triggers calcium release from the sarcoplasm (By similarity). Plays an important role in excitation-contraction coupling in the heart. Required for normal heart development and normal regulation of heart rhythm (PubMed:21216955). Required for normal contraction of smooth muscle cells in blood vessels and in the intestine. Essential for normal blood pressure regulation via its role in the contraction of arterial smooth muscle cells (PubMed:14609949, PubMed:28119464). Long-lasting (L-type) calcium channels belong to the 'high-voltage activated' (HVA) group (Probable).[UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (9) differs in the 5' UTR and has multiple coding region differences, compared to variant 1. These differences cause translation initiation at an alternate AUG and result in an isoform (9) that is shorter and has a distinct N-terminus, compared to isoform 1. 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.