

Product datasheet for SC301591

CREB5 (NM_001011666) Human Untagged Clone

Product data:

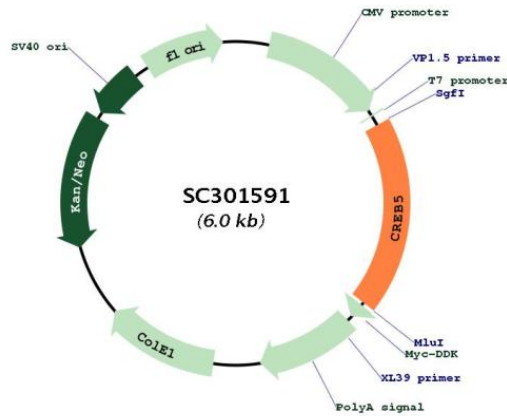
| | |
|---------------------------|--|
| Product Type: | Expression Plasmids |
| Product Name: | CREB5 (NM_001011666) Human Untagged Clone |
| Tag: | Tag Free |
| Symbol: | CREB5 |
| Synonyms: | CRE-BPA; CREB-5; CREBPA |
| Mammalian Cell Selection: | Neomycin |
| Vector: | pCMV6-Entry (PS100001) |
| E. coli Selection: | Kanamycin (25 ug/mL) |
| Fully Sequenced ORF: | >SC301591 representing NM_001011666. Blue=Insert sequence Red=Cloning site Green=Tag(s) |

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GCTCGTTTAGTGAACCGTCAGAATTTGTAAACGACTCACTATAGGGCGCCGGGAATTCGTCGACTG
GATCCGGTACCGAGGAGATCTGCCGCCGCGATCGCC
ATGTTCTGCACCTCAGGAGGGAATTCAGCCTCAGTGATGTCCATGAGGCCTGTCCCAGGCTCTCTATCT
TCTCTGTACATCTCCACAACAGACAGAGACAGCCCATGCCAGCCTCCATGCCTGGGACCCTGCCAAC
CCTACAATGCCAGGATCTCCGCCGTCTTGATGCCAATGGAGCGACAAATGTCAGTGAATCCAGCATC
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GCCAAAATGAGGTTGAAGGCTGCATTGACTCACCACCCTGCTGCCATGTCAAATGGGAACATGAACACC
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CCGCATCAGCACCAGACTGCCACCCATCACCTTACCACACCAGCACCAGCACCAGCACACCAT
CCTCACCCCTCAACCCATCACCAGCAGAACCATCCACATCACCCTCCCATTCCCACCTTCATGCACAC
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ACACAACAGATGCAGCAACCCAGACAATACAGCCACCCAGCCACAGGGGGCGCCGGCGAAGGGTG
GTAGACGAGGATCCGGACGAGAGGCGCGGAAATTTCTGGAACGGAACCGGGCAGCTGCCACCCGCTGC
AGACAGAAGAGGAAGGTCTGGGTGATGTCATTGAAAAAGAAAGCAGAAGAACTCACCCAGACAAACATG
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GCTAGTCCTGTCCAGTTGCTCCAGCAACAAGTCATCCAGCATAATACCATCACTACTTCCTCATCG
GTCAGCGAGGTGGTAGGAAGCTCCACCCTCAGCCAGCTCACCCTCACAGAACAGACCTGAATCCGATT
CTTTAA
ACGCGTACGCGGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAATGATATCCTGGAT
TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC
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Restriction Sites: SgfI-MluI



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Plasmid Map:


ACCN: NM_001011666

Insert Size: 1110 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: This TrueClone is provided through our Custom Cloning Process that includes sub-cloning into OriGene's pCMV6 vector and full sequencing to provide a non-variant match to the expected reference without frameshifts, and is delivered as lyophilized plasmid DNA.

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_001011666.2](#)

RefSeq Size: 7917 bp

RefSeq ORF: 1110 bp

Locus ID: 9586

UniProt ID: [Q02930](#)

Cytogenetics: 7p15.1
Protein Families: Transcription Factors
Protein Pathways: Huntington's disease, Prostate cancer

MW: 41.2 kDa

Gene Summary: The product of this gene belongs to the CRE (cAMP response element)-binding protein family. Members of this family contain zinc-finger and bZIP DNA-binding domains. The encoded protein specifically binds to CRE as a homodimer or a heterodimer with c-Jun or CRE-BP1, and functions as a CRE-dependent trans-activator. Alternatively spliced transcript variants encoding different isoforms have been identified. [provided by RefSeq, Jul 2008]
Transcript Variant: This variant (4) lacks several 5' exons and uses an alternate 5' terminal exon which results in the use of a downstream start codon, compared to variant 1. It encodes isoform delta which is shorter and contains a distinct N-terminus compared to isoform alpha.
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.