

Product datasheet for SC109101

CREM (NM_182720) Human Untagged Clone

Product data:

OriGene Technologies, Inc.

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Product Type:	Expression Plasmids
Product Name:	CREM (NM_182720) Human Untagged Clone
Tag:	Tag Free
Symbol:	CREM
Synonyms:	CREM-2; hCREM-2; ICER
Mammalian Cell Selection:	None
Vector:	pCMV6-XL4
E. coli Selection:	Ampicillin (100 ug/mL)
Fully Sequenced ORF:	>OriGene ORF within SC109101 sequence for NM_182720 edited (data generated by NextGen Sequencing) ATGGCTGTAACTGGAGATGACACAGCTGCCACTGGTGACATGCCAACTTACCAGATCCGA GCTCCTACTGCTGCTTTGCCACAGGGAGTGGTGATGGCTGCATCGCCCGGAAGTTTGCAC AGTCCCCAGCAGCTGGCAGAAGAAGCAACACGCAAACGAAGGCTGAGGCTAATGAAAAAC AGGGAAGCTGCCCGGGAGTGTCGCAGGAAGAAGAAAGAATATGTCAAATGTCTTGAAAAT CGTGTGGCTGTGCCTGAAAACCAAAACAAGACTCTCATTGAGGAACTCAAGGCCCTCAAA GATCTTTATTGCCATAAAGTAGAGTAA

Clone variation with respect to NM_182720.1



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5' Read Nucleotide Sequence:	<pre>>OriGene 5' read for NM_182720 unedited GGATTTTGTAATACGACTCACTATAGGGCGGCCGCGCAATTCGGCACGAGGGNAAACAANA CAGTTCTGTCTGCAGAAGCCCATTATGGCTGTAACTGGAGATGACACAGCTGCCACTGGT GACATGCCAACTTACCAGATCCGAGCTCCTACTGCTGCTTTGCCACAGGGAGTGGTGATG GCTGCATCGCCCGGAAGTTTGCACAGTCCCCAGCAGCTGGCAGAAGAAGCAACACGCAAA CGAGAGCTGAGGCTAATGAAAAACAGGGAAGCTGCCCGGGAGTGTCGCAGGAAGAAGAAA GAATATGTCAAATGTCTTGAAAATCGTGTGGCTGTGCTTGAAAACCAAAACAAGACTCTC ATTGAGGAACTCAAGGCCCTCAAAGATCTTTATTGCCATAAAGTAGAGTAACTGTCTTTG ACTTGGAGCTTGTTTACTCTAATCAAGGCAGGAGGAGGAGCAGCAGCAGCTCTACTTATTGCCATG TGGACTTGTGGGAAGGACACGTGTGACCCTTAAGAATCCAGTTTGGATTAGTGTTTGAAA TTGAATTGGGAATGTTGTTCCAGGATGTGGAATGCAGCCGTGATCACACTTACTGACAT ACTTTGAGGAAGGACACGTGTGGAATGCAGCCGTGAACAACAAGAACTTGTCTGCTT TTTTTCAGGGAAGCTGCCAAAGAATGCCGCGTGAACAAGAAAGA</pre>
Restriction Sites:	Notl-Notl
ACCN:	NM_182720
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:	 Centrifuge at 5,000xg for 5min. Carefully open the tube and add 100ul of sterile water to dissolve the DNA. Close the tube and incubate for 10 minutes at room temperature. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom. 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:	<u>NM 182720.1, NP 874389.1</u>
RefSeq Size:	1955 bp
RefSeq ORF:	327 bp
Locus ID:	1390
UniProt ID:	<u>Q03060</u>
Cytogenetics:	10p11.21
Protein Families:	Druggable Genome, Transcription Factors

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Gene Summary:

This gene encodes a bZIP transcription factor that binds to the cAMP responsive element found in many viral and cellular promoters. It is an important component of cAMP-mediated signal transduction during the spermatogenetic cycle, as well as other complex processes. Alternative promoter and translation initiation site usage allows this gene to exert spatial and temporal specificity to cAMP responsiveness. Multiple alternatively spliced transcript variants encoding several different isoforms have been found for this gene, with some of them functioning as activators and some as repressors of transcription. [provided by RefSeq, Jul 2008]

Transcript Variant: This variant (7), also known as ICER1 gamma and hCREM 2alpha-b, uses an alternative downstream promoter, and differs in the 5' UTR, 3' UTR, and coding region, compared to variant 1. This results in a shorter isoform (7, also known as g) with distinct N-and C-termini, compared to isoform 1. This isoform represents one of the early repressor ICER isoforms. 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.

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