

## Product datasheet for RN216427

### Cebpa (NM\_001287578) Rat Untagged Clone

#### Product data:

**Product Type:** Expression Plasmids  
**Product Name:** Cebpa (NM\_001287578) Rat Untagged Clone  
**Tag:** Tag Free  
**Symbol:** Cebpa  
**Synonyms:** DBPCEP  
**Vector:** pCMV6-Entry (PS100001)  
**E. coli Selection:** Kanamycin (25 ug/mL)  
**Cell Selection:** Neomycin  
**Fully Sequenced ORF:** >RN216427 representing NM\_001287578  
Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC  
GCCGGATCGC

ATGAGCAGCCACCTCCAGAGCCCCCGCAGCGCCAGCAACGCCGCCTTTGGCTTTCCCGGGGCGCGG  
GCCCGCGCCGCCAGCCCCACCTGCCGCCCGGAGCCGCTGGCGGCATCTGCGAGCAGGACGTC  
TATAGACATCAGCGCCTACATCGACCCGCGCCCTTCAACGACGAGTTCTGGCCGACCTTTCCAGCAC  
AGCCGGCAGCAGGAGAAGGCCAAGGCGCGCGGGCCCCGCGGGTGGCGCGGTGACTTTGACTACCCGG  
GCGCCCCGCGGGCCCCGGCGGTGCGGTATGTCCGCGGGGGCGCACGGACCCCCCTCCGGCTACGGCTG  
TGCGGGCGCCGGCTACCTGGACGGCAGGCTGGAGCCCCGTACGAGCGCGTGGGGCGCCCGCGCTGCGG  
CCGCTGGTGATCAAGCAGGAGCCCCGCGAGGAGGACGAGGCGAAGCAGCTGGCGCTGGCCGGCCTTTCC  
CCTATCAGCCCCGCGCCGCCGCCACCACCGCACGCGTCTCCCGCGCACTTGGCCGCCCTCA  
CTTGCAAGTTCCAGATCGCACACTGCGGCCAGACCACCATGCACCTGCAGCCTGGCCACCTACGCCGCC  
CCGACGCCCGTGCCAGCCCTCATCCCGCGCTGCAATGGTGCTGCGGGCTGCCGGCCCCGGGGCGC  
CGCTCAAGGGCTTGGCTGGTCCGACCCCGACCTCCGCACCGCGCGGGCGCGGGCGGGGGCGCGC  
GGGCAAGGCCAAGAAGTCGGTGGATAAGAAGCAGCAACGAGTACCGGGTACGGCGGGAACGCAACAACATC  
GCGGTGCGCAAGACCGAGATAAAGCCAAACAGCGCAACGTGGAGACGACGAGAGGTGTTGGAGTTGA  
CCAGTGACAATGACCGCTGCGCAAGCGGTGGAACAGCTGAGCCGTGAACTGGACACGCTGCGGGTAT  
CTTCCGCCAGCTGCTGAGAGCTCCTTGGTCAAGGCCATGGGCAACTGCGCGTGA

AGCGGACCGACGCTACGCGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCC  
TGGATTACAAGGATGACGACGATAAGGTTTAA

**Restriction Sites:** SgfI-RsrII  
**ACCN:** NM\_001287578



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<b>Insert Size:</b>	1035 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>OTI Annotation:</b>	Clone contains native stop codon, and expresses the complete ORF without any c-terminal tag.
<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_001287578.1</a></u> , <u><a href="#">NP_001274507.1</a></u>
<b>RefSeq Size:</b>	2673 bp
<b>RefSeq ORF:</b>	1035 bp
<b>Locus ID:</b>	24252
<b>UniProt ID:</b>	<u><a href="#">P05554</a></u>
<b>Cytogenetics:</b>	1q21
<b>Gene Summary:</b>	<p>This intronless gene encodes a transcription factor that contains a basic leucine zipper (bZIP) domain and recognizes the CCAAT motif in the promoters of target genes. The encoded protein functions in homodimers and also heterodimers with CCAAT/enhancer-binding proteins beta and gamma. Activity of this protein can modulate the expression of genes involved in cell cycle regulation as well as in body weight homeostasis. The use of alternative in-frame non-AUG (CUG) and AUG start codons results in protein isoforms with different lengths. Differential translation initiation is mediated by an out-of-frame, upstream open reading frame which is located between the CUG and the first AUG start codons. [provided by RefSeq, Aug 2014]</p> <p>Transcript Variant: This variant (1) can initiate translation from an upstream non-AUG (CUG) site, and also from three conserved downstream, in-frame AUG sites. The isoform (d) represented in this RefSeq results from translation initiation at the second AUG start codon. Isoform d has a shorter N-terminus, compared to isoform c. 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>