

## Product datasheet for **MC227657**

### Ctbp1 (NM\_001198860) Mouse Untagged Clone

#### Product data:

**Product Type:** Expression Plasmids  
**Product Name:** Ctbp1 (NM\_001198860) Mouse Untagged Clone  
**Tag:** Tag Free  
**Symbol:** Ctbp1  
**Synonyms:** BARS; CtBP3/BARS; D4S115h; D5H4S115; D5H4S115E  
**Vector:** pCMV6-Entry (PS100001)  
**E. coli Selection:** Kanamycin (25 ug/mL)  
**Cell Selection:** Neomycin  
**Fully Sequenced ORF:** >MC227657 representing NM\_001198860  
 Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC  
 GCC**GCGATCGCC**

ATGTCAGGCGTCCGACCTCCCATCATGAACGGGCCATGCACCCCGGCCCTGGTGGCGCTGCTGGATG  
 GCCGGGACTGCACAGTGGAGATGCCTATCCTGAAGGATGTGGCCACAGTAGCCTTCTGTGATGCACAGTC  
 CACACAGGAGATCCATGAGAAGTACTGAATGAGGCTGTGGGTGCCCTGATGTACCATACCATCACACTG  
 ACCAGAGAAGATCTGGAGAAGTTAAAGCTCTTAGAATCATCGTCCGAATTGGCAGCGGGTTTGACAATA  
 TCGACATCAAGTCAGCTGGGGATCTAGGCATCGCAGTGTGCAATGTGCCGCGAGCATCTGTGGAAGAAAC  
 GGCAGACTCCACCTGTGCCACATCCTGAACCTGTACCGACGAACCACCTGGCTACACCAGGCCCTTCGG  
 GAAGGCACTCGAGTCCAGAGTGTAGAGCAGATCCGAGAGGTGGCTTCAGGAGCTGCCAGGATCCGTGGAG  
 AGACCTTGGGCATCATTGGACTAGGTCTGTGGCCAGGCGTGGCACTTCGGGCAAAGGCTTTTGGCTT  
 CAACGCTCTTCTATGATCCATACCTATCTGATGGAATCGAGCGGGCCCTGGGGCTACAGCGCGTGAGC  
 ACGTGCAGGACCTGCTCTTCCACAGTACTGCGTTACCCTGCATTGCGGCCTCAATGAGCACAACCACC  
 ACCTCATCAATGACTTTACTGTCAAGCAGATGAGACAAGGAGCCTTTCTGGTGAACACAGCCCGTGGTGG  
 CCTGGTGGATGAGAAGGCACTGGCCAGGCCCTGAAGGAAGGGCGGATCCGTGGCGCAGCGCTGGACGTG  
 CATGAGTCAGAGCCCTTACGCTTTAGCCAGGGACCCTAAAGGATGCACCAACCTCATCTGCACACCCC  
 ATGCTGCATGGTACAGTGAAGGCGTCCATTGAGATGAGAGAGGAGGCAGCCCGGAAATCCGGCGAGC  
 CATCACAGGCCGATCCCAGATAGCTTGAAAACTGTGTCAACAAGGACCACCTGACAGCCGCCACGCAC  
 TGGGCCAGCATGGACCCTGCTGTGGTGCACCCTGAGCTCAATGGGGTGCCTACAGGTACCTCCAGGCG  
 TCGTGAGTGTGGCCCCACTGGCATCCCAGCTGCTGTGGAAGGGATTGTTCCAGTGCCATGTCCCTGTC  
 TCATGGCCTGCCCTGTGGCCACCCACCCACGCTCCCTCTCCTGGCCAGACTGTCAAGCCTGAGGCG  
 GATAGAGACCATACGAGTGACCAGTT**AG**

**ACGCGT**ACGCGGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT  
 ACAAGGATGACGACGATAAGGTTTAA



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<b>Restriction Sites:</b>	Sgfl-Mlul
<b>ACCN:</b>	NM_001198860
<b>Insert Size:</b>	1290 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_001198860.1</a></u> , <u><a href="#">NP_001185789.1</a></u>
<b>RefSeq Size:</b>	2495 bp
<b>RefSeq ORF:</b>	1290 bp
<b>Locus ID:</b>	13016
<b>UniProt ID:</b>	<u><a href="#">O88712</a></u>
<b>Cytogenetics:</b>	5 17.52 cM
<b>Gene Summary:</b>	<p>Corepressor targeting diverse transcription regulators such as GLIS2 or BCL6. Has dehydrogenase activity. Involved in controlling the equilibrium between tubular and stacked structures in the Golgi complex. Functions in brown adipose tissue (BAT) differentiation. [UniProtKB/Swiss-Prot Function]</p> <p>Transcript Variant: This variant (2) contains an alternate exon in the 5' region, initiates translation at an alternate start codon, and uses an alternate in-frame splice site in the 3' coding region, compared to variant 3. It encodes a shorter isoform (2) with a distinct N-terminus compared to isoform 3.</p>