

## Product datasheet for **MC201337**

### **Tfcp2 (NM\_033476) Mouse Untagged Clone**

#### **Product data:**

Product Type:	Expression Plasmids
Product Name:	Tfcp2 (NM_033476) Mouse Untagged Clone
Tag:	Tag Free
Symbol:	Tfcp2
Synonyms:	CP-2; CP2; D230015P20Rik; LBP-1c; LBP-1d; LBP1; LSF; Tcfcp2; UBP-1
Mammalian Cell Selection:	Neomycin
Vector:	PCMV6-Kan/Neo (PCMV6KN)
E. coli Selection:	Kanamycin (25 ug/mL)



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**Fully Sequenced ORF:** >BC022131 sequence for NM\_033476  
 CTTTTCGCTGTCCACGCTTCCAGGCGGTCTGGTGGCTAAGGAGGGTGTGGGGACGGCTCGGAGGAACCCCT  
 GGGAGCCGGCGCCTGTTGGGGCAAGGAAGGAGCCGGATGGCCTGGGCTCTGAAGCTGCCGCTAGCCGAC  
 GAAGTGATCGAGTCGGGGCTTGTGCAGGACTTTGATGCTAGCCTGTCCGGCATTGGCCAGGAACTGGGTG  
 CTGGTGCTTATAGTATGAGTGATGTCCTCGCATTGCCATTTTTAAGCAAGAAGAGTCAAGTTTGCCTCC  
 CGATAATGAGAACGAGATTCTGCCTTTTCAGTATGTGCTCTGTGCTGCCACGCTCCACGGGTGAAGCTC  
 CACGACGAGACCCTGACGTACCTCAATCAAGGACAGTCTTATGAAATCCGAATGCTAGACAATAGAAAA  
 TGGGAGAGCTTCCAGAACTCAACGGCAAGCTGGTGAAGAGCATATTCCGGTAGTGTCCACGACAGGCG  
 ACTGCAGTACACTGAGCACCAGCAGCTGGAGGGTGGAGGTGGAACCGACCGGGAGACAGGATTCTCGAC  
 ATAGATATCCCAATGTCTGTGGGTGTAATTGATCCCAGGGCCAATCCTACCCAGCTAAACACAGTGGAGT  
 TCCTGTGGGACCCTTCAAAGAGGACATCTGTGTTTATTAGGTGCACTGCATCAGCACTGAGTTCACCAT  
 GAGGAAGCATGGCGGGGAGAAGGGGTGCCGTTCCGAGTGCAGATCGATACCTTCAAGGAGAACGGGAAC  
 GGAGAGTACACAGAGCACTTGCCTCGCCAGCTGCCAGATCAAAGTCTTCAAGCCTAAGGGTGCAGACA  
 GAAAACAAAAAATAGATAGAGAGAAAATGGAGAAACGGACGCCCATGAAAAAGAGAAAATCAACCTTC  
 TTATGAGACAACCATACTCACCGAGTGTCTCCGTGGCCTGAGATCACCTACGTCAACAATTCCCATCC  
 CCTGGCTTCAACAGCTCCCATAGCAGTTTCTCTTGGGGAAGGAAACGGCTCGCCCAACCACCAGCCAG  
 AACCCACCCCTCCAGTACCGGATAACCTCTTGGCGACAACCACGCCTCAGGAAGCCACGCAATGGCTGCA  
 TCGAAACCGGTTCTCCACATTCACGAGGCTTTTACCAACTTCTCAGATTTACTGAAACTAACTAGAGAC  
 GACGTGATCCAAATCTGCGGCCCTGCAGATGGAATCAGACTCTTAAATGCATTAAGGCGCCGATGGTGC  
 GGCCAAGGCTAACATTTATGTCTGCCAGGAGTCGTTGCAGTTGAGGGGAGCAGCAGCCGACGCCAGCC  
 GCAGCCGAGAGCAGGAGGATGGGGATCAAACGGTACTTTCTCGTGTACCATGCTATCTATCTCGAG  
 GAGTGCAGCTGTGCAACTCACAGAAAAATGCTCAGCTTTTTCAGATTTCTCCTCACCAGATCAGCC  
 AGATTTACAAGCAAGGGCCGACAGGGATCCATGTGGTCATCAGCGATGAGATGGTACAGAACTCCAGGA  
 AGAAGCCTGCTTTATTCTGGACACGATGGAAGCAGAAACAGTACAGCTATCATGTCTCAAGTAG  
 GGTGAGGCCCGACCTTCGACGGCTGCTGCTGCCCTTCCCTCTTACCTTCTGAAGGGGACATGACTGGA  
 GATTAAGGGCTTCAGGAACCTGGCACATCTGAATGTTGGGGGAGGGGGAGGCGTGTCCAGTGCAGGG  
 CGAGCAGGTCCCAAGGAGGAATCCAGAGTCTGTATTGGCTCCTCTGATCACTGGCTATACATGGAGCC  
 AGCAGTTTCTTACTTTCTGGCCCTGTTGATACCAGTAAGAAGCCCGTCATTAGGGTTGTAGAGGGC  
 AGAGTGGTAATGGGAGTCATGTGATGTTTTCTAGGAAGATGTTTAGTTTTGTGATATAGAGCTCTGACAT  
 TTTACTGATATAAAAGTGAACATCTGTCTTCATAAAACCAAGCTCTCTCTTGCCTCAAGCCTGTTGGC  
 AGCTCTGCTCTGCTCTGTTCCAACTGTGAGGAGGAGAAACAGAACTCTCTCCTTCCCTTTTTTGGAGACG  
 GGTTCATGTTCTTGCCTTACCTCCCAAGAGCTGGGATCACAAGCTTTTCCACCTGGTCTGTTTCTGT  
 CTGCAAGTCTGGGCTTGAACCTCGGAGCCTCACGTGCTAGGCAGGCACTTTCTCCAGCTGAGCTCCAGCAG  
 GCCTTCCCTTTAGTGCTAAGACAGGCTCCAGATTTCTAATTTGGTTACCTGCAATACCACAGCTGAGTG  
 TGATTTTGGCTAATGTGTATTGGGGTAGACCGAATGATAAATCCCACGGACGGATAATCTTCTTTTATCC  
 CCTTAATAAAACCAAAACATCTGGGGACACAGTGAAGGTTGAGGGGAGGAGACGCAGATGAAGTGGAGGGG  
 CTTCTGACAACACCAGATGCCACCCTTGTGCTGCAAGTGTCCACTGGGCTTTTCTGGGGCTCTGCTGG  
 AAAGTACGAGCACTTCCACTGAAGGAGGCAACCATACCTTACCAGTGTAGCGATAATGAAGCCCAAGACA  
 GACAGACACCAATGTCACACTAACCTCAGCCAGAGCTAGAGGTGAGAGAGACAAGCGTGTCTGACCC  
 AGGCTTCGTTGTTGCGTAATGATATTCTCAAACTGTTTAAAGTAAATGAGCTATTCGCGCCTGCTCTCCC  
 CTCCCCTGCTTGTGTTTGGCATAGTGTGGTGTGTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTAAATGTAAG  
 CTTTATTAATAAAAAAGAAAAATACATTCTTTTCTTCTTCAAACTCACACTGTGTTTATGTTCTGCCTTTCA  
 AATGGCTGTGTCTGTGTGCATGTACAGAATTGTAAGCACTTAGGCCCTCCTTTTTTACATGAAGTTT  
 GTTTTAGCTTGGTTTTCCAAGAAAACGTTTCTCTGTGGAGCCCGGCTGTCTGGAACCTCACTCTGTAGA  
 CCAGGCTGGCCTCAAACCTCAGAAATCCACCTTCCAAGTGTGGGAATATAGGTGTGTGCCACTAGCTTAG  
 TCAGGCCACATGTAAGTTTTACCAGAAATCCCTTAAATTTTTGTTGAGATGAACAAAGATAATGTTAAA  
 GATTCTGTATTAATTTGAATTTATGTGTGTTTTGAAGTATTTAATAAAGCTTAGATTTTTTTATAGAG  
 AAAAAAAAAAAAAAAAAA

**Restriction Sites:** RsrII-NotI  
**ACCN:** NM\_033476

<b>Insert Size:</b>	1503 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>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="#">BC022131</a></u> , <u><a href="#">AAH22131</a></u>
<b>RefSeq Size:</b>	3238 bp
<b>RefSeq ORF:</b>	1503 bp
<b>Locus ID:</b>	21422
<b>UniProt ID:</b>	<u><a href="#">Q9ERA0</a></u>
<b>Cytogenetics:</b>	15 56.33 cM
<b>Gene Summary:</b>	<p>Binds a variety of cellular promoters including fibrinogen, alpha-globin promoters. Activation of the alpha-globin promoter in erythroid cells is via synergistic interaction with UBP1 (By similarity). Functions as part of the SSP (stage selector protein) complex. Facilitates the interaction of the gamma-globin genes with enhancer elements contained in the locus control region in fetal erythroid cells. Interacts by binding to the stage selector element (SSE) in the proximal gamma-globin promoter (By similarity).[UniProtKB/Swiss-Prot Function]</p> <p>Transcript Variant: This variant (1) represents the longer transcript and encodes the longer 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.</p>