

## Product datasheet for **MC226568**

### **Bdnf (NM\_001285418) Mouse Untagged Clone**

#### Product data:

**Product Type:** Expression Plasmids  
**Product Name:** Bdnf (NM\_001285418) Mouse Untagged Clone  
**Tag:** Tag Free  
**Symbol:** Bdnf  
**Vector:** pCMV6-Entry (PS100001)  
**E. coli Selection:** Kanamycin (25 ug/mL)  
**Cell Selection:** Neomycin  
**Fully Sequenced ORF:** >MC226568 representing NM\_001285418  
Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC  
GCC**CGATCGC**C

ATGACCATCCTTTTCCTTACTATGGTTATTTCACTTTCGGTTGCATGAAGGCGGCGCCCATGAAAGAAG  
TAAACGTCACGACAAGGCAACTTGGCCTACCCAGGTGTGCGGACCCATGGGACTCTGGAGAGCGTGAA  
TGGGCCAGGGCAGGTTTCGAGAGGTCTGACGACGACATCACTGGCTGACACTTTTGAGCAGTCATCGAA  
GAGCTGCTGGATGAGGACCAGAAGTTTCGGCCCAACGAAGAAAACCATAAGGACGCGGACTTGTACTACT  
CCCGGGTGATGCTCAGCAGTCAAGTGCCTTTGGAGCCTCCTACTCTTTCTGCTGGAGGAATACAAAA  
TTACCTGGATGCCGAAACATGTCTATGAGGGTTCGGCGCCACTCCGACCCTGCCCGCCGTGGGAGCTG  
AGCGTGTGTGACAGTATTAGCGAGTGGGTACAGCGGCAGATAAAAAGACTGCAGTGGACATGTCTGGCG  
GGACGGTCACAGTCCTAGAGAAAGTCCCGGTATCCAAAGGCCAACTGAAGCAGTATTTCTACGAGACCAA  
GTGTAATCCCATGGGTTACACCAAGGAAGGCTGCAGGGGCATAGACAAAAGGCACTGGAACCTCGCAATGC  
CGAACTACCAATCGTATGTTTCGGGCCCTTACTATGGATAGCAAAAAGAGAATTGGCTGGCGATTATAA  
GGATAGACTTCTGTGTATGTACTGACCATTAAGGGGAAGTAG

**ACGCGT**ACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT  
ACAAGGATGACGACGATAAGGTTTAA

**Restriction Sites:** SgfI-MluI  
**ACCN:** NM\_001285418  
**Insert Size:** 750 bp



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<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_001285418.1</a></u> , <u><a href="#">NP_001272347.1</a></u>
<b>RefSeq Size:</b>	3866 bp
<b>RefSeq ORF:</b>	750 bp
<b>Locus ID:</b>	12064
<b>UniProt ID:</b>	<u><a href="#">P21237</a></u>
<b>Cytogenetics:</b>	2 56.63 cM
<b>Gene Summary:</b>	<p>The protein encoded by this gene is a member of the nerve growth factor family. It is involved in the growth, differentiation and survival of specific types of developing neurons both in the central nervous system (CNS) and the peripheral nervous system. It is also involved in regulating synaptic plasticity in the CNS. Expression of a similar gene in human is reduced in both Alzheimer's and Huntington disease patients. Alternative splicing results in multiple transcript variants encoding different isoforms that may undergo similar processing to generate mature protein. [provided by RefSeq, Oct 2015]</p> <p>Transcript Variant: This variant (7) differs in the 5' UTR and coding region, and uses a downstream translation start compared to variant 1. The resulting protein (isoform 2) has a shorter N-terminus compared to isoform 1. Variants 2, 3, 4, 5, 6, 7, 8, 9, 10, 11 and 12 encode the same protein. 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>