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# Product datasheet for SC328480

## MBNL3 (NM_001170702) Human Untagged Clone

### Product data:

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| **Product Type:** | Expression Plasmids |
| **Product Name:** | MBNL3 (NM_001170702) Human Untagged Clone |
| **Tag:** | Tag Free |
| **Symbol:** | MBNL3 |
| **Synonyms:** | CHCR; MBLX; MBLX39; MBXL |
| **Mammalian Cell Selection:** | Neomycin |
| **Vector:** | pCMV6-Entry (PS100001) |
| **E. coli Selection:** | Kanamycin (25 ug/mL) |
| **Fully Sequenced ORF:** | >SC328480 representing NM_001170702.<br>Blue=Insert sequence Red=Cloning site Green=Tag(s)<br><br>GCTCGTTTAGTGAACCGTCAGAATTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTG<br>GATCCGGTACCGAGGAGATCTGCCGCCGCGATCGCC<br>ATGGAGAGGGCTTCTAAGAATCTTAAGGGTCGGTGTACCCGAGAGAACTGCAAGTACCTTCACCCTCCT<br>CCACACTTAAAAACGCAGCTGGAGATTAATGGGCGGAACAATCTGATTCAACAGAAGACTGCCGCAGCC<br>ATGTTCGCCCAGCAGATGCAGCTTATGCTCCAAAACGCTCAAATGTCATCACTTGGTTCTTTTCCTATG<br>ACTCCATCAATTCCAGCTAATCCTCCCATGGCTTTCAATCCTTACATACCACATCCTGGGATGGGCCTC<br>GTTCCTGCAGAACTTGTACCAAATACACCTGTTCTGATTCCTGGAAACCCACCTCTTGCAATGCCAGGA<br>GCTGTTGGCCCAAAACTGATGCGTTCAGATAAACTGGAGGTTTGCCGAGAATTTCAGCGTGGAAATTGT<br>ACCCGTGGGGAGAATGATTGCCGCTATGCTCACCCTACTGATGCTTCCATGATTGAAGCGAGTGATAAT<br>ACTGTGACAATCTGCATGGATTACATCAAAGGTCGATGCTCGCGGGAGAAATGCAAGTACTTTCATCCT<br>CCTGCACACTTGCAAGCCAGACTCAAGGCAGCTCATCATCAGATGAACCATTCAGCTGCCTCTGCCATG<br>GCCCTGCAGCCTGGTACACTGCAACTGATACCAAAGAGATCAGCACTGGAAAAGCCCAATGGTGCCACC<br>CCGGTCTTTAATCCCACTGTTTTCCACTGCCAACAGGCTCTGACTAACCTGCAGCTCCCACAGCCGGCA<br>TTTATCCCTGCAGTGCCCATGATGCACGGTGCTACACCTACCACTGTGTCTGCAGCAACAACACCTGCC<br>ACCAGCGTTCCGTTCGCTGCACCAACTACAGGCAATCAGCTGAAATTCTGA<br>ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGAT<br>TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC |
| **Restriction Sites:** | SgfI-MluI |

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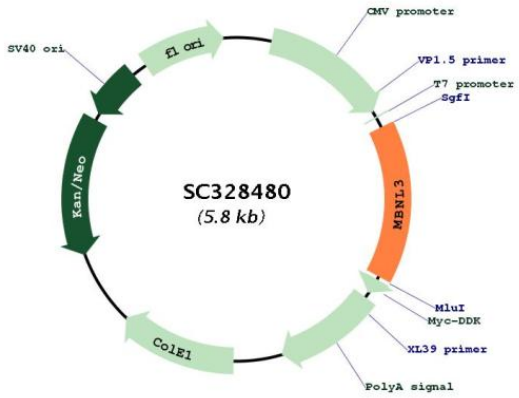
Plasmid Map:

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| **ACCN:** | NM_001170702 |
| **Insert Size:** | 879 bp |
| **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). |
| **OTI Annotation:** | This TrueClone is provided through our Custom Cloning Process that includes sub-cloning into OriGene's pCMV6 vector and full sequencing to provide a non-variant match to the expected reference without frameshifts, and is delivered as lyophilized plasmid DNA. |
| **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:** | 1. Centrifuge at 5,000xg for 5min.<br>2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.<br>3. Close the tube and incubate for 10 minutes at room temperature.<br>4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.<br>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. |
| **RefSeq:** | NM_001170702.1 |
| **RefSeq Size:** | 11303 bp |
| **RefSeq ORF:** | 879 bp |

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| **Locus ID:** | 55796 |
| **UniProt ID:** | Q9NUK0 |
| **Cytogenetics:** | Xq26.2 |
| **MW:** | 31.8 kDa |
| **Gene Summary:** | This gene encodes a member of the muscleblind-like family of proteins. The encoded protein may function in regulation of alternative splicing and may play a role in the pathophysiology of myotonic dystrophy. Alternatively spliced transcript variants have been described. [provided by RefSeq, Dec 2009]<br>Transcript Variant: This variant (4) differs in the 5' UTR, lacks a portion of the 5' coding region, initiates translation at an alternate start codon, and lacks an in-frame exon in the 3' coding region compared to variant 1. The encoded isoform (4) has a distinct N-terminus and is shorter than 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. |

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