

OriGene Technologies, Inc.

9620 Medical Center Drive, Ste 200 Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com CN: techsupport@origene.cn

Product datasheet for SC315963

Dystrophia myotonica protein kinase (DMPK) (NM_001081560) Human Untagged Clone

Product data:

| Product Type: | Expression Plasmids |
|------------------------------|--|
| Product Name: | Dystrophia myotonica protein kinase (DMPK) (NM_001081560) Human Untagged Clone |
| Tag: | Tag Free |
| Symbol: | DMPK |
| Synonyms: | DM; DM1; DM1PK; DMK; MDPK; MT-PK |
| Mammalian Cell Selection: | Neomycin |
| Vector: | pCMV6-Entry (PS100001) |
| E. coli Selection: | Kanamycin (25 ug/mL) |
| | |



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| |)ystrophia myotonica protein kinase (DMPK) (NM_001081560) Human Untagged Clone – C315963 |
|---------------------------|--|
| Fully Sequenced (| <pre>RF: >SC315963 representing NM_001081560. Blue=Insert sequence Red=Cloning site Green=Tag(s)</pre> |
| | GCTCGTTTAGTGAACCGTCAGAATTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTG GATCCGGTACCGAGGAGATCTGCCGCC <mark>GCGATCGCC</mark> |
| | ATGTCAGCCGAGGTGCGGCTGAGGCGGCTCCAGCAGCTGGTGTTGGACCCGGGCTTCCTGGGGCTGGAG |
| | CCCCTGCTCGACCTTCTCCTGGGCGTCCACCAGGAGCTGGGCGCCTCCGAACTGGCCCCAGGACAAGTAC |
| | GTGGCCGACTTCTTGCAGTGGGCGGAGCCCATCGTGGTGAGGCTTAAGGAGGTCCGACTGCAGAGGGAC GACTTCGAGATTCTGAAGGTGATCGGACGCGGGGGCGTTCAGCGAGGTAGCGGTAGTGAAGATGAAGCAG |
| | ACGGGCCAGGTGTATGCCATGAAGATCATGAACAAGTGGGACATGCTGAAGAGGGGGCGAGGTGCGTGC |
| | TTCCGTGAGGAGAGGGACGTGTTGGTGAATGGGGACCGGCGGTGGATCACGCAGCTGCACTTCGCCTTC |
| | CAGGATGAGAACTACCTGTACCTGGTCATGGAGTATTACGTGGGCGGGGACCTGCTGACACTGCTGAGC |
| | AAGTTTGGGGAGCGGATTCCGGCCGAGATGGCGCGCTTCTACCTGGCGGAGATTGTCATGGCCATAGAC |
| | TCGGTGCACCGGCTTGGCTACGTGCACAGGGACATCAAACCCGACAACATCCTGCTGGACCGCTGTGGC |
| | CACATCCGCCTGGCCGACTTCGGCTCTTGCCTCAAGCTGCGGGCAGATGGAACGGTGCGGTCGCTGGTG |
| | GCTGTGGGCACCCCAGACTACCTGTCCCCCGAGATCCTGCAGGCTGTGGGCGGTGGGCCTGGGACAGGC AGCTACGGGCCCGAGTGTGACTGGTGGGCGCCTGGGTGTATTCGCCTATGAAATGTTCTATGGGCAGACG |
| | CCCTTCTACGCGGATTCCACGGCGGAGACCTATGGCAAGATCGTCCACTACAAGGAGCACCTCTCTG |
| | CCGCTGGTGGACGAAGGGGTCCCTGAGGAGGCTCGAGACTTCATTCA |
| | ACACGGCTGGGCCGGGGTGGAGCAGGCGACTTCCGGACACATCCCTTCTTCTTTGGCCTCGACTGGGAT |
| | GGTCTCCGGGACAGCGTGCCCCCCTTTACACCGGATTTCGAAGGTGCCACCGACACATGCAACTTCGAC |
| | TTGGTGGAGGACGGGCTCACTGCCATGGAGACACTGTCGGACATTCGGGAAGGTGCGCCGCTAGGGGTC |
| | CACCTGCCTTTTGTGGGCTACTCCTACTCCTGCATGGCCCTCAGGGACAGTGAGGTCCCAGGCCCCACA |
| | CCCATGGAACTGGAGGCCGAGCAGCTGCTTGAGCCACACGTGCAAGCGCCCAGCCTGGAGCCCTCGGTG |
| | TCCCCACAGGATGAAACAGCTGAAGTGGCAGTTCCAGCGGCTGTCCCTGCGGCAGAGGCTGAGGCCGAG GTGACGCTGCGGGAGCTCCAGGAAGCCCTGGAGGAGGAGGTGCTCACCCGGCAGAGCCTGAGCCGGGAG |
| | ATGGAGGCCATCCGCACGGACAACCAGAACTTCGCCAGTCAACTACGCGAGGCAGAGGCTCGGAACCGG |
| | GACCTAGAGGCACACGTCCGGCAGTTGCAGGAGCGGATGGAGTTGCTGCAGGCAG |
| | GTCACGGGGGTCCCCAGTCCCCGGGCCACGGATCCACCTTCCCATCTAGATGGCCCCCCGGCCGTGGCT |
| | GTGGGCCAGTGCCCGCTGGTGGGGCCAGGCCCCATGCACCGCCGCCACCTGCTGCCCGGGGCCAGGGTC |
| | CCTAGGCCTGGCCTATCGGAGGCGCTTTCCCTGCTCCTGTTCGCCGTTGTTCTGTCTCGTGCCGCC |
| | ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGAT |
| | TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC |
| Restriction Sites: | Sgfl-Mlul |
| Plasmid Map: | |
| ACCN: | NM_001081560 |
| Insert Size: | 1875 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. |

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| Dystrophia myotonica protein kinase (DMPK) (NM_001081560) Human Untagged Clone – SC315963 | |
|---|--|
| 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 Met | hod: 1. Centrifuge at 5,000xg for 5min. 2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA. 3. Close the tube and incubate for 10 minutes at room temperature. 4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom. 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: | <u>NM 001081560.2</u> |
| RefSeq Size: | 2859 bp |
| RefSeq ORF: | 1875 bp |
| Locus ID: | 1760 |
| UniProt ID: | <u>Q09013</u> |
| Cytogenetics: | 19q13.32 |
| Protein Families: | Druggable Genome, Protein Kinase |
| MW: | 69 kDa |
| Gene Summary: | The protein encoded by this gene is a serine-threonine kinase that is closely related to other kinases that interact with members of the Rho family of small GTPases. Substrates for this enzyme include myogenin, the beta-subunit of the L-type calcium channels, and phospholemman. The 3' untranslated region of this gene contains 5-38 copies of a CTG trinucleotide repeat. Expansion of this unstable motif to 50-5,000 copies causes myotonic dystrophy type I, which increases in severity with increasing repeat element copy number. Repeat expansion is associated with condensation of local chromatin structure that disrupts the expression of genes in this region. Several alternatively spliced transcript variants of this gene have been described, but the full-length nature of some of these variants has not been determined. [provided by RefSeq, Jul 2016] Transcript Variant: This variant (3) has multiple differences in the presence and absence of exons at its 5' end and in the CDS, compared to variant 1. These differences produce a distinct 5' UTR, cause translation initiation at an alternative start codon, and the loss of an in-frame portion of the coding region, compared to variant 1. The encoded protein (isoform 3, also known as isoform 11) 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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