

## Product datasheet for **MC222906**

### Piwil2 (NM\_021308) Mouse Untagged Clone

#### Product data:

Product Type:	Expression Plasmids
Product Name:	Piwil2 (NM_021308) Mouse Untagged Clone
Tag:	Tag Free
Symbol:	Piwil2
Synonyms:	mili; Piwil11
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
Cell Selection:	Neomycin



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Fully Sequenced ORF:

>MC222906 representing NM\_021308

Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC  
GCC**CGCATCGCC**

ATGGATCCTGTACAGGCCGTTGTTTCAGGGGGCCACCCAGTCCACCCATCTCAGTGTGTGGGATGCCAG  
GCTGTTGGCCTCAAGCTCCTAGACCTTTGGAACCAAGCTTGGGGTAGGGCAGGACCTGCAGGCAGAGGCCT  
TGTGTTTAGAAAACAGAACTCCAGCCACCACTCCAGCCAGTACAAAAGATTCTGTAGGTTTGGTG  
TCCATGTTCCGTGGCATGGGTCTTGACACAGCATTCCGGCCTCCTTCAAAACGAGAAGTGCCCTCTTTAG  
GCAGAGGAGTTCTAGGTCGAGGCTTGTCTGCTAACATGGTCCGCAAGGACAGAGAAGAACCCGTTCCCTC  
TTTGCCTGATCCTTCGGTGTGGCAGCTGGGGACAGCAAAGTGGCAGAGGCTTCTGTTGGTTGGAGTAGA  
ATGCTGGGAAGAGGTAGTTCTGAGGTCTCTGTTACCACTGGGACGAGCAGCCAGTATAGGCAGAG  
GAATGGACAAACCTCCCAGTGCCTTTGGCCTCACAGCTCGGGATCCCCACGGCTGCCACAGCCTCCAGC  
TCTGTCTCCAACCTCACTGCACTCTGCCGATCCCCTCCGGTCTGACTATGGAACGAAAGGAAAAAGAG  
CTTTTGGTCAAGCAAGGATCAAAAGGAACCTCTCAGTCTTTGGGACTGAACCTCATCAAAATCCAGTGTC  
ATAACGAAGCAGTTTATCAGTACCATGTGACTTTCAGCCCCAGTGTGGAATGCAAAAGCATGAGGTTTGG  
CATGTTGAAGGACCACAGTCTGTCACTGGAAACGTCACTGCTTTTATGGCTCTATTCTTTATCTTCTCT  
GTTAAGCTTCAACAAGTTGTTGAGTTAAAAAGTCAGAGGAAAAGTACGATGCCGAGATCAGTATCAAGA  
TTCAGTGCACAAAGATCCTGGAGCCGTTTCTGACCTGTGCATCCCCTTCTACAATGTTGTCTTCCGGCG  
GGTAATGAAACTTCTGGATATGAAGCTTGTGGGGAGAACTTCTATGACCCTACAAGTCCATGGTACTG  
CAGCAACACAGATTGCAGATCTGGCCTGGCTATCGGGCTAGTATCCGGAGGACAGACGGGGTCTCTTCC  
TGCTCGTGTATCTCTCATAAGGTCACTCGGAACGACTCTGTGCTGGATGTCATGCATGCTATCTACCA  
GCAGAACAAGGAGCACTTCCAGGACGAGTGCAGCAAGCTTCTGGTTGGCAGCATTGTATCACGCGCTAC  
AACAATCGTACCTACCGAATCGATGTGGACTGGAACAAGACCCCTAAAGACAGCTTTGTATGTCTCGG  
ACGGGAAGGAAATCACATTCTGGAATACTACAGCAAAAAGTATGGGATCACAGTCAAGGAAGATGACCA  
GCCGCTGTGATCCACCGGCCAGTGAGAGACAGAATAACCATGGCATGTTGCTGAAGGGCAGATCCTG  
CTGCTGCCGAGCTCTCTTTCATGACGGGGATCCCTGAGAAGTGAAGAAGGACTTCAGGGCCATGAAGG  
ACTTGACTCAGCAGATTAACCTGAGCCCCAAGCAGCACCACGGTCTTTGGAATGCCTGTCAGAGAAT  
TTCACAAAACGAGACAGCCAGCAATGAGCTGACCCGCTGGGGCTCAGTCTGCATAAAGATGTCCACAAG  
ATTGAAGGTGGCTTCTGCCAATGGAGAGGATCAACTTAAGGAACACTTCATTTGTCACATCGGAGGACC  
TGAAGTGGTTAAGGAAGTGACCAGAGATGCTTCCATTCTAACTATTCCCATGCATTTCTGGGCACTCTT  
TTATCAAAGAGAGCAATGGACCAAGCCAGAGAAGTGGTTAACATGTTGAAAAGATTGCCGGGCCATT  
GGCATGCCATAAGCCCCCAGCCTGGGTTGAGCTGAAGGATGACCGAATAGAGACCTATATCAGGACCA  
TTCAGTCTTACTGGGAGTTGAGGGGAAGATACAAATGGTCGTTTGCATCATATGGGCACACGTGATGA  
TCTCTATGGAGCCATCAAGAAGCTGTGCTGCGTGCAGTCCCAGTGCCCTCACAGGTATCAATGTCCGA  
ACCATTGGTCAGCCACCAGGCTTCGGAGCGTGGCTCAGAAAATTTTACTTCAGATGAACTGAACTGG  
GTGGTGAGCTCTGGGAGTGGATATTCCGCTGAAACAATAATGGTGATTGGAATGGATGTGTACCATGA  
CCCCAGCAGAGGCATGCGCTCTGTGGTGGCTTCGTGGCCAGCATAAATCTCACACTACCAAAATGGTAC  
TCGAGGGTGGTGTCCAGATGCCACATCAGGAGATTGTGGACAGCCTGAAGCTCTGCCTGGTGGTTCTCT  
TGAAAAAGTATTATGAGGTGAACCATTTGCTCCCAGAGAAAATTGTGGTGTACCGAGATGGAGTGTCTGA  
TGGCCAGCTAAAGACAGTTGCCAACTACAGATCCCTCAGCTGCAGAAGTGTGTTGAAGCCTTTGATAAC  
TACCACCCCAAGATGGTGGTGTGTTGAGTTTCAAGAAAATCAGCACCAATCTGTACCTTGTCTGCTCCTG  
ATCACTTCTGAACCCCTCCCCGGGACTGTGGTTGATCATACCATAACCAGCTGTGAGTGGGTGGATTT  
CTACCTTCTTGGCCATCATGTGCGACAGGGCTGTGGCATACCTACACACTACATCTGTGTTCTGAACACT  
GCAAACTGAGCCCTGATCATGCAGAGGTTGACTTTCAAATATGCCACATGACTGGAATTGGCCTG  
GTACCATCCGAGTTCAGCTCCTTGAAGTATGCCACAAGCTAGCTTTCCTGTCCGGACAGATTTTGCA  
TCATGAGCCAGCCATCCAGCTGTGTGGAACTGTTCTTCTGTAA

**ACGCGT**ACGCGGCCGCTCGAGCAGAAAAGTCACTCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT  
ACAAGGATGACGACGATAAGGTTTAA

<b>Restriction Sites:</b>	Sgfl-Mlul
<b>ACCN:</b>	NM_021308
<b>Insert Size:</b>	2916 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="#">NM_021308.1</a></u> , <u><a href="#">NP_067283.1</a></u>
<b>RefSeq Size:</b>	4913 bp
<b>RefSeq ORF:</b>	2916 bp
<b>Locus ID:</b>	57746
<b>UniProt ID:</b>	<u><a href="#">Q8CDG1</a></u>
<b>Cytogenetics:</b>	14 D2

**Gene Summary:**

Endoribonuclease that plays a central role during spermatogenesis by repressing transposable elements and preventing their mobilization, which is essential for the germline integrity (PubMed:11578866, PubMed:14736746, PubMed:17446352, PubMed:18381894, PubMed:18922463, PubMed:26669262). Plays an essential role in meiotic differentiation of spermatocytes, germ cell differentiation and in self-renewal of spermatogonial stem cells (PubMed:11578866, PubMed:14736746, PubMed:17446352, PubMed:18381894, PubMed:18922463, PubMed:26669262). Its presence in oocytes suggests that it may participate in similar functions during oogenesis in females (PubMed:11578866, PubMed:14736746, PubMed:17446352, PubMed:18381894, PubMed:18922463, PubMed:26669262). Acts via the piRNA metabolic process, which mediates the repression of transposable elements during meiosis by forming complexes composed of piRNAs and Piwi proteins and govern the methylation and subsequent repression of transposons (PubMed:11578866, PubMed:14736746, PubMed:17446352, PubMed:18381894, PubMed:18922463, PubMed:26669262). During piRNA biosynthesis, plays a key role in the piRNA amplification loop, also named ping-pong amplification cycle, by acting as a 'slicer-competent' piRNA endoribonuclease that cleaves primary piRNAs, which are then loaded onto 'slicer-incompetent' PIWIL4 (PubMed:22020280, PubMed:23706823, PubMed:26669262). PIWIL2 slicing produces a pre-miRNA intermediate, which is then processed in mature piRNAs, and as well as a 16 nucleotide by-product that is degraded (PubMed:28633017). Required for PIWIL4/MIWI2 nuclear localization and association with secondary piRNAs antisense (PubMed:18381894, PubMed:18922463, PubMed:26669262). Besides their function in transposable elements repression, piRNAs are probably involved in other processes during meiosis such as translation regulation (PubMed:19114715). Indirectly modulates expression of genes such as PDGFRB, SLC2A1, ITGA6, GJA7, THY1, CD9 and STRA8 (PubMed:16261612). Represses circadian rhythms by promoting the stability and activity of core clock components ARNTL/BMAL1 and CLOCK by inhibiting GSK3B-mediated phosphorylation and ubiquitination-dependent degradation of these proteins (PubMed:28903391).[UniProtKB/Swiss-Prot Function]