

## Product datasheet for MC227348

### Hormad1 (NM\_001289537) Mouse Untagged Clone

#### Product data:

**Product Type:** Expression Plasmids  
**Product Name:** Hormad1 (NM\_001289537) Mouse Untagged Clone  
**Tag:** Tag Free  
**Symbol:** Hormad1  
**Synonyms:** 4921522K05Rik; Nohma  
**Vector:** pCMV6-Entry (PS100001)  
**E. coli Selection:** Kanamycin (25 ug/mL)  
**Cell Selection:** Neomycin  
**Fully Sequenced ORF:** >MC227348 representing NM\_001289537  
 Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC  
 GCC**GCGATCGC**C

ATGGCCACTATGCAGTTGCAGAGGACAGCTTCCCTGAGTGCATTGGTATTTCCAATAAGATATCAACTG  
 AGCATCAATCTTTGATGTTTGTGAAGAGGCTCCTAGCTGTTTCAGTATCTTGCATCACCTATTTGAGAGG  
 AATATTTCCAGAACGTGCTTATGGGACAAGATATCTGGATGATCTCTGTGTCAAAATCTGAAAGAAGAT  
 AAAAAATTGTCCAGTTCTTACAGCTAGTGAAGTGGATGCTTGGATGCTATGATGCTTTACAGAAGAAAT  
 ATCTAAGGATGATCATTCTAGCTGTATACACCAATCCAGGAGATCCTCAGACAATTTGAGAAATGTTACCA  
 GTTTAAATTAAGTACACCAAAAATGGACCAATCATGGACTTTATAAGCAAAAATCAAAAACAATAAATCT  
 AGTACAACATCTGCTGACACCAAGAAAGCAAGTATTCTCCTCATTGGAAGATTTATGTCTTAATGCAAA  
 ATCTAGGACCATACCTAATGATGTTTGTCTGACCATGAACTTTTTACTATGATGAAGTTACACCCCC  
 AGATTACCAACCACCAGGTTTTAAGGATGGTACTGTGAAGGAGTAATATTTGATGGGGACCCTACATAC  
 TTAATGTGGGAGAAGTCCCAACACCTTTTACACCTTCAGATTAAGGAGTACTGAGAAAGGAAAGGAA  
 TGGAAAAATATTGATCAACCATACTAAAACAAAAGAATCAAAAACAAATTTGAAAAAATCTAATGGA  
 CAAAGATGATGTGAAGATGAAAAATCATAATAATTTGACATTAAGGAGTAAAGTAAAGGAAAGGAAAGT  
 AACTCTGGAGCTTCTGAAATCAAAGAACCAAAATTTAGATTGTAAGGAAAGAAAGAACTATGCAATTCAAA  
 AGAGCCAAAGTCTTCAATTTCTCATTGTGAGGTTGAACAGTTAGTCAGTAAAACATCTGAACCTTGATG  
 GTCTGAAAGCAAAAACAAGAAGCGGAAAAATCTTTCAGAGTAAAATGGTAAATGAAATAATCAACAAGGA  
 CAAACTTCTAAGAAAATCGGAAGAGAAGTCTTCGTCAATTTAGGAAAACAATAAATGCACCTGAGTGTA  
 GGTGA

**ACGCGT**ACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT  
 ACAAGGATGACGACGATAAGGTTTAA

**Restriction Sites:** SgfI-MluI



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<b>ACCN:</b>	NM_001289537
<b>Insert Size:</b>	1125 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>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_001289537.1</a></u> , <u><a href="#">NP_001276466.1</a></u>
<b>RefSeq Size:</b>	1438 bp
<b>RefSeq ORF:</b>	1125 bp
<b>Locus ID:</b>	67981
<b>UniProt ID:</b>	<u><a href="#">Q9D5T7</a></u>
<b>Cytogenetics:</b>	3 F2.1
<b>Gene Summary:</b>	Plays a key role in meiotic progression (PubMed:19686734, PubMed:21079677, PubMed:21478856). Regulates 3 different functions during meiosis: ensures that sufficient numbers of processed DNA double-strand breaks (DSBs) are available for successful homology search by increasing the steady-state numbers of single-stranded DSB ends (PubMed:19686734, PubMed:21079677). Promotes synaptonemal-complex formation independently of its role in homology search (PubMed:19686734, PubMed:21079677). Plays a key role in the male mid-pachytene checkpoint and the female meiotic prophase checkpoint: required for efficient build-up of ATR activity on unsynapsed chromosome regions, a process believed to form the basis of meiotic silencing of unsynapsed chromatin (MSUC) and meiotic prophase quality control in both sexes (PubMed:21478856).[UniProtKB/Swiss-Prot Function] Transcript Variant: This variant (3) differs in the 3' UTR and coding sequence compared to variant 1. The resulting isoform (b) has a shorter and distinct C-terminus compared to isoform a. Variants 2, 3, and 4 all encode the same isoform (b).