

Product datasheet for **MC224345**

Nsd2 (NM_175231) Mouse Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	Nsd2 (NM_175231) Mouse Untagged Clone
Tag:	Tag Free
Symbol:	Nsd2
Synonyms:	5830445G22Rik; 9430010A17Rik; AW555663; C130020C13Rik; D030027O06Rik; D930023B08Rik; mKIAA1090
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
Cell Selection:	Neomycin
Fully Sequenced ORF:	>MC224345 representing NM_175231 Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
GCC**CGATCGCC**

ATGGAATTTAGCATCAGAAAAAGTCCTCTTTCTGTTTCAGAAAGTTGTAAGTGTCATGAAGATGAAGCAGA
CACCAGAAATCCTTGGCAGTGCAAATGGAAAGACTCAGAACTGTGAAGTGAATCATGAATGTTCTGTATT
CCTCAGCAAAGCTCAACTTTCTAACAGCCTACAGGAGGGGTCATGCAGAAATTTAATGGCCATGATGCC
CTCCCCCTTTCTCCAGCAGAGAAGTTGAAAGATCTTACTTCTTGTGTTTTAATGGAGAACCTGGTGCTC
ATGATACTAAATTGTGTTTTGAGGCCAGGAAGTAAAAGGAATTGGACACCACCAATACTACACCTAT
CAAAAATGGCTCTCCAGAAATTAACCTGAAAATACCAAAACATACATGAATGGGAAACCTCTTTTGAA
TCTTCAATTTGTGGTGACGGTGCTGATGTGTCTCAGTCAGAAGAAAACGAACAAAACTGATAATA
AGACTAGGAGGAACAGGAAGAGGAGCATAAAAATGACTCTTTACTGGAGCAGGGCCTTGTGGAAGCAGC
TTAGTGTCCAAGATCTCAAGTCTGCAGATAAAAAGATTCCAGTTAAGAAGGAGTCTGTCCAAACTACT
GGCAGAGACAGAGACCTTTGTTAAAATACAACGTTGGTGATTTGGTGTGGTCCAAGTGTGAGTTACC
CTTGGTGGCCTTGCATGGTTTCTGCTGACCCACTCCTTCAACATCATACCAAACTTAAAGGTCAGAAAA
AAGTGCACGCCAGTATCATGTACAGTCTTTGGTGATGCCCCAGAAAGAGCTTGGATATTTGAGAAGAGC
AAGCTGAGAAAAATCAAGTTGTTGAAACCTATTTGGGGAGATTGAGAGCCAGTGGGAAATGGGCATTGT
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TGGACTCTTCAGGGGCCAGTGAAGAGGCTGCTGTAGACCCTGGGTCTGTGAGAGAAGAGGATATCCAC
GAAGAGAAGGCCAAGAACCAAAGGTCTAGTCTGCTGAGAACCAAGAAGGTGATCTGGCACAGACAAG
AGTACACCTCAAAGATGGCAGAGGCTGAACCTAAGCGAGGAGTAGGCTCTCTGCTGGGAGGAAAAAGT
CCACAGGCTCTGCTCCTCGGAGCAGGAAGGAGACTCAGCAGCCAGTTTTTAGTCTTTTGTCAAAAACA
CAGAGATGAGTTGTAGCTGAACATCCAGATGCCTCAGGGGAAGAGATTGAAGAATTGCTTGGTCCCAG



TGGAGCATGCTCAATGAAAAGCAGAAAGCAGCATATAATACAAAGTTTTCCCTAATGATCTCTGCCAGT
 CTGAAGAAGACTCTGAAATGGGAATGGGAAAAAAGAAGCCACACAAAGAGAGCAGATGACCCTGCAGA
 GGATGTGGATGTTGAAGACGCGCCAGGAAAAGACTTAGAGCAGATAAGCACAGTCTTCGGAAGAGAGAG
 ACAATCACCGACAAAACGGCCAGAAACAGCTCTTACAAGGCTATAGAGGCAGCCTCCTCAAGAGCC
 AGGCAGCAACGAAAAATTTGTCTGATGCTTGCAAACCACTGAAGAAGCGAAATCGGGCTTCTGCAACAGC
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 AGCCCAGGAGATGAACCATCAGAGTCTCCATATGAAAGTGCAGATGAAACACAGACTGAAGCATCCGTCT
 CATCTAAAAAATCTGAAAGAGGAATGGCTGCCAAAAAGGAGTACGTGTGTGAGTGTGAGAAAGACAGG
 CAGTCTCTTACTGTGTGAGGGCCCTGTTGTGGAGCATTCCACCTAGCCTGCCTTGGACTTCCCGAAGA
 CCGGAAGGAAGATTACCTGCACCGAATGTGCCTCAGGGATTCACTCATGCTTCGTATGTAAGAAAGCA
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 CCCTCTCACTGTGTTGAGAGCCGAGGCTTCGCTGTCTCTCCATAGCTGTATGAGCTGCCATGCCTCC
 AACCTTCAAATCCAGGCCATCAAAGGTAAATGATGCGATGTGTCCGATGCCCGTTGCCTATCATG
 GAGGGGATGCCTGTCTGGCAGCAGGATGTTGCGTGATTGCTTCTAACAGCATAATCTGCACAGGCCACTT
 CACTGCCCGAAGGGGAAGCGGCACCATACCCATGTCAATGTGAGCTGGTCTTTGTGTGCTCAAAGGG
 GGAAGCCTTCTGTGCTGTGAGGCCTGCCAGCAGCCTTTCACCCGATTGTCTAAACATCGAGATGCCCG
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 ACTTGGTAACTACAGATGGTGGCCGGCAGAAAGTTGCCATCCAAAAATGTTCCCCCAAATATTCAGAAA
 ATGAAGCACGAGATTGGAGAATCCCTGTATTTTTCTTTGGGTCTAAAGATTATTACTGGACGCATCAGG
 CACGAGTGTCCCATACATGGAGGGGACCGGGCAGCCGCTACCAGGGGTGAGAGGATCGGAAGAGT
 CTTCAAGAACGCATTGCAAGAAGCTGAAGCTCGTTTTAATGAAGTCAAACCTCAGCGGAAGCCCGAAG
 ACACAGGAGAGTGAGAGAAAGCCTCCACCATAACAAGCATCAAGGTGAATAAACCTTATGGTAAAGTCC
 AGATCTACACAGCTGACATTTCTGAAATCCCAGTGAAGTCAAGTCAAGTCAAGTCAAGTCAAGTCAAGT
 CTCTGATTGAGAGTGTCTGAACAGGATGCTAATGTTTGAAGTGCCACCCGAGGTGTCTGCAGGGGAA
 TACTGCCAGAACCAGTGTTCCTAAGCCAGTACCCTGAGACCAAGATCATCAAGACAGATGGCAAAG
 GGTGGGCCTGTTGCCAAAAGGACATCAGAAAAGGAGAATTTGTTAATGAGTATGTTGGTGGAGCTGAT
 TGATGAAGAGGAGTGTATGGCAAGAATCAAATATGCACATGAGAATGACATCACTCACTTCTACATGCTC
 ACCATAGACAAGGACCGCATAATTGATGCTGGCCCAAAGGGAATTATTCAGATTATGAACCATAGCT
 GCCAGCCCAATTGTGAGACCCTGAAGTGGACAGTGAATGGGACACACGGGTTGGCCTGTTTGTGTGTG
 TGACATTCCTGCAGGTACAGAGCTGACTTCAACTATAACCTTGATTGTTTGGGCAATGAAAAGACAGTC
 TGTCGGTGTGGAGCCTCAACTGTAGTGGTTCCTTGGAGACAGACCAAGACATCAGCATCCCTTTTCAT
 CAGAGGAAAAGGGTAAAAAGGCCAAGAAGAAAACAGAAAGGCGCAGAGCCAAGGGTGAAGGAAAACGGCA
 GTCAGAGGATGAGTGTCCGCTGTGGTGTGGTGGGCAGCTGGTGTGTGTGACCGCAAGTTCTGTACC
 AAGGCCTACCACCTATCCTGCCTCGGTTTGGGCAAGCGGCCCTTTGGGAAGTGGGAATGTCTTGGCATC
 ACTGTGATGATGTGGCAAACCTTCTACCTCATTTTGGCACCTCTGCCCAAACCTATTCTGTAAGGAAACA
 CCAAGATGGGACTGCTTCCGTTCCACCAGGATGGGCAGTCTACTGCTGTGAGCATGACTTGAGGGCA
 GACTCTTCAAGTAGCACCAAGACTGAGAAACCCCTTCCAGAATCACTGAAGTCAAAGGAAAAGAGGAAGA
 AAAGGCGGTGCTGGCGAAGGGTCACAGATGGCAAA**TAG**

ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT
 ACAAGGATGACGACGATAAGGTTTAA

Restriction Sites:

SgfI-MluI

ACCN:

NM_175231

Insert Size:

4098 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).

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:	<ol style="list-style-type: none">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_175231.2</u> , <u>NP_780440.2</u>
RefSeq Size:	6933 bp
RefSeq ORF:	4098 bp
Locus ID:	107823
UniProt ID:	<u>Q8BVE8</u>
Cytogenetics:	5 B2
Gene Summary:	<p>Histone methyltransferase with histone H3 'Lys-27' (H3K27me) methyltransferase activity. [UniProtKB/Swiss-Prot Function]</p> <p>Transcript Variant: This variant (2) uses an alternate in-frame splice site in the central coding region, compared to variant 1, resulting in an isoform (2) that is 1 aa shorter than isoform 1. The 5' UTR of this variant may be incomplete due to the lack of 5'-complete transcripts supporting it, and the presence of alternative splicing choices further upstream. 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>