

## Product datasheet for **SC317003**

### HDAC2 (NM\_001527) Human Untagged Clone

#### Product data:

Product Type:	Expression Plasmids
Product Name:	HDAC2 (NM_001527) Human Untagged Clone
Tag:	Tag Free
Symbol:	HDAC2
Synonyms:	HD2; KDAC2; RPD3; YAF1
Mammalian Cell Selection:	None
Vector:	<u><a href="#">pCMV6-XL5</a></u>
E. coli Selection:	Ampicillin (100 ug/mL)



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**Fully Sequenced ORF:** >OriGene ORF sequence for NM\_001527 edited  
 CAGCAGCAGCAGCAGCAGCAGCAGCAGGAGGAGCCCGGTGGCGGCGGTGGCCGGGAGCC  
 CATGGCGTACAGTCAAGGAGGCGCAAAAAAAAAAGTCTGCTACTACTACGACGGTGATAT  
 TGGAAATTATTATTATGGACAGGGTCATCCCATGAAGCCTCATAGAATCCGCATGACCCA  
 TAACCTGCTGTTAAATTATGGCTTATACAGAAAAATGAAATATATAGGCCCATAAAGC  
 CACTGCCGAAGAAATGACAAAATATCACAGTGATGAGTATATCAAATTTCTACGGTCAAT  
 AAGACCAGATAACATGTCTGAGTATAGTAAGCAGATGCAGAGATTTAATGTTGGAGAAGA  
 TTGTCCAGTGTTTGTGGACTCTTTGAGTTTTGTGAGCTCTCAACTGGCGGTTTCAGTTGC  
 TGGAGCTGTGAAGTTAAACCGACAACAGACTGATATGGCTGTTAATTGGGCTGGAGGATT  
 ACATCATGCTAAGAAATCAGAAGCATCAGGATTCTGTTACGTTAATGATATTGTGCTTGC  
 CATCCTTGAATTACTAAAGTATCATCAGAGAGTCTTATATATTGATATAGATATTCATCA  
 TGGTGATGGTGTGAAGAAGCTTTTTATAACAACAGATCGTGAATGACGGTATCATTCCA  
 TAAATATGGGGAATACTTTCCTGGCACAGGAGACTTGAGGGATATTGGTGTGGAAAAGG  
 CAAATACTATGCTGTCAATTTTCCAATGAGAGATGGTATAGATGATGAGTCATATGGGCA  
 GATATTTAAGCCTATTATCTCAAAGGTGATGGAGATGTATCAACCTAGTGTGTGGTATT  
 ACAGTGTGGTGCAGACTCATTATCTGGTATAGACTGGGTGTTTCAATCTAACAGTCAA  
 AGGTCATGCTAAATGTGTAGAAGTTGTA AAAA ACTTTTAACTTACCATTACTGATGCTTGG  
 AGGAGGTGGCTACACAATCCGTAATGTTGCTCGATGTTGGACATATGAGACTGCAGTTGC  
 CCTTGATTGTGAGATTCCCAATGAGTTGCCATATAATGATTACTTTGAGTATTTTGGACC  
 AGACTTCAAACATGCATATTAGTCCTTCAAACATGACAAACCAGAACACTCCAGAATATAT  
 GGAAAAGATAAAACAGCGTTTGTGTTGAAAATTTGCGCATGTTACCTCATGCACCTGGTGT  
 CCAGATGCAAGCTATCCAGAAGATGCTGTTTATGAAGACAGTGGAGATGAAGATGGAGA  
 AGATCCAGACAAGAGAATTTCTATTCGAGCATCAGACAAGCGGATAGCTTGTGATGAAGA  
 ATTCTCAGATTCTGAGGATGAAGGAGAAGGAGGTGCAAGAAATGTGGCTGATCATAAGAA  
 AGGAGCAAAGAAAGCTAGAATTGAAGAAGATAAGAAAAGAAAACAGAGGACAAAAAACAGA  
 CGTTAAGGAAGAAGATAATCCAAGGACAACAGTGGTGA AAAA ACAGATACCAAAGGAAC  
 CAAATCAGAACAGCTCAGCAACCCTGAATTTGACAGTCTACCAATTTTCAAGAAATCAT  
 TAAAAAGAAAATATTGAAAGGAAAATGTTTTCTTTTTGAAGACTTCTGGCTTCATTTTAT  
 ACTACTTTGGCATGGACTGTATTTATTTTCAAATGGCTTTTTCTGTTTTGTTTTCTTGG  
 CAAGTTTTATTGTGAGTTTTTCTAATTATGAAGCAAAAATTTCTTTTCTCCACCATGCTTT  
 ATGTGATAGTATTTAAAATTGATGTGAGTATTATGTCAAAAAACTGATCTATTAAGA  
 AGTAATTGGCCTTCTGAGCTGATTTTTCCATCTTTTGAATTATCTTTATTA AAAA AT  
 GTACTTGGAAAAAAA

**5' Read Nucleotide Sequence:** >OriGene 5' read for NM\_001527 unedited  
 TACCTTGTATACGACTCCTATAGGGCGGCCGGAATTCGGCACGAGGATGCGCTCACCTC  
 CCTGCGGCCTCCTGAGGTGGTTTTGGTGGCCCCCTCCTCGCGAGTTGGTGCCGCTGCCACC  
 TCCGATTCGAGCTTTCGGCACCTCTGCCGGGTGGTACCAGCCTTCCCGCGCCCCCTC  
 CTCTCCTCCCACCGCCTGCCCTTCCCCGCGGACTATCGCCCCACGTTTCCCTCAGCC  
 CTTTTCTCTCCCGCCGAGCCGCGGCGGCAGCAGCAGCAGCAGCAGCAGCAGCAGGAGGA  
 GGAGCCCCGTGGCGGCGGTGGCCGGGAGCCCATGGCGTACAGTCAAGGAGGCGGCAAAA  
 AAAAAGTCTGCTACTACTACGACGGTGATATTGAAAATTATTATTATGGACAGGGTCATC  
 CCATGAAGCCTCATAAGATCCGCATGACCATAACTTGTCTGTTAAATTATGGCTTATACA  
 GAAAAATGAAAATATATAGGCCCATAAAGCCACTGCCGAAGAAATGACAAAATATCACA  
 GTGATGAGTATATCAAATTTCTACGGTCAATAAGACCAGATAACATGTCTGAGTATAGTA  
 AGCAGATGCAGAGATTTAATGTTGGAGAAGATTGTCCAGTGTGTTGATGGACTCTTTGAGT  
 TTTGTGAGCTCTCAACTGGCGGTTTCAGTTGCTGGAGCTGTGAAGTTAAACCCGACAACAG  
 ACTGATATGGCTTGTAAATTTGGCTGGAGATTACATCATGCTAAGAAATCAGAGCATCAG  
 ATTCTGTTACGTAATGATATTTGTGCTGCATCTGATACTAAGTATCATCAGAAGAGTCTA  
 TATATTGAATATAGATATCATCATGTGATTGTGTTGAGAGCTTTTTATACAAACAGATCG  
 GTGATGACGTATTCATTCCT

<b>3' Read Nucleotide Sequence:</b>	<p>&gt;OriGene 3' read for NM_001527 unedited            CGAGATGCACTTCAGGGCCGGAAGCACCGGGTGAGGGTCACAGGGATGCCACCCGGGAT            CTGTTTCAGGAAACAGCTATGACCGCGGCCCAATCTAGAGTCGAGTTTTTTTTTTTTTTT            TTTTCCAAGTACAATTTTTTAATAAAGATAATTACAAAAGATGGAAAAATCAGCTCAAAA            AGGCCAATTACTTCTTTAATAGATCAGTTTTTTTGACATAATAACTCACATCAATTTTAA            ATACTATCACATAAAGCATGGTGGAAAAAATTTTGCTTCATAATTAGAAAACTCA            CAATAAACTTGCCAAGAAAAACAAAAACGAAAAAGCCATTTGAAAATAAATACAGTCCA            TGCCAAAAGTAGTATAAAATGAAGCCAGAAGTCTTCAAAAAGAAAACATTTTCCTTTCAAT            ATTTTCTTTTTAATGATTTTCTGAAATTGGTGAGACTGTCAAATTCAGGGTTGCTGAGC            TGTCTGATTTGGTTCCTTTGGTATCTGTTTTTTCACCACTGTTGTCCTTGGATTATCT            TCTTCTTAACGTCTGTTTTTTGTCCTCTGTTTCTTTCTTATCTTCTTCAATTCTAGCT            TTCTTTGCTCCTTTCTTATGATCAGCCACATTTCTTCGACCTCCTTCTCCTTCATCCTCA            GAATCTGAGAATTCTTCATCACAAGCTATCCGCTTGTCTGATGCTCGAATAGAAATTCTC            TTGTCTGGATCTTCTCCATCTTCATCTCCACTGTCTTCATGACAGCATCTTCTGGAATAG            CTTGCATCTGGACACAGTGCATGAGTACATGCGCAATTTTCAAACAACGCTGTTTATCTT            TCATATATTCTGGAGTGTCTGTTGTCATGTGAGACTATATGCAGTGAGTGTCCAATACT            CAGTATCATATGCACTCATGGCATCTCCATCAGGCACTGCG</p>
<b>Restriction Sites:</b>	Please inquire
<b>ACCN:</b>	NM_001527
<b>Insert Size:</b>	1900 bp
<b>OTI Disclaimer:</b>	<p>Due to the inherent nature of this plasmid, standard methods to replicate additional amounts of DNA in E. coli are highly likely to result in mutations and/or rearrangements. Therefore, OriGene does not guarantee the capability to replicate this plasmid DNA. Additional amounts of DNA can be purchased from OriGene with batch-specific, full-sequence verification at a reduced cost. Please contact our customer care team at <a href="mailto:custsupport@origene.com">custsupport@origene.com</a> or by calling 301.340.3188 option 3 for pricing and delivery.</p> <p>The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing variants is recommended prior to use. <a href="#">More info</a></p>
<b>OTI Annotation:</b>	<p>This clone may be unstable or toxic at high copy number in common E. coli strain. We recommend using a lower copy number E. coli strain, such as CopyCutter strain (<a href="http://www.epibio.com/item.asp?ID=435">http://www.epibio.com/item.asp?ID=435</a>) for transformation and plasmid preparation. Please be aware that the DNA yield could be low. Additional aliquots of this clone can be ordered from OriGene.</p>
<b>Components:</b>	<p>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).</p>

**Reconstitution Method:**

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:** [NM\\_001527.2](#), [NP\\_001518.2](#)

**RefSeq Size:** 6659 bp

**RefSeq ORF:** 1749 bp

**Locus ID:** 3066

**UniProt ID:** [Q92769](#)

**Cytogenetics:** 6q21

**Domains:** Hist\_deacetyl

**Protein Families:** Druggable Genome, Stem cell - Pluripotency, Transcription Factors

**Protein Pathways:** Cell cycle, Chronic myeloid leukemia, Huntington's disease, Notch signaling pathway, Pathways in cancer

**Gene Summary:** This gene product belongs to the histone deacetylase family. Histone deacetylases act via the formation of large multiprotein complexes, and are responsible for the deacetylation of lysine residues at the N-terminal regions of core histones (H2A, H2B, H3 and H4). This protein forms transcriptional repressor complexes by associating with many different proteins, including YY1, a mammalian zinc-finger transcription factor. Thus, it plays an important role in transcriptional regulation, cell cycle progression and developmental events. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Apr 2010]  
Transcript Variant: This variant (1) encodes the functional protein. 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.