

Product datasheet for **SC331503**

HDAC9 (NM_001204144) Human Untagged Clone

Product data:

Product Type: Expression Plasmids
Product Name: HDAC9 (NM_001204144) Human Untagged Clone
Tag: Tag Free
Symbol: HDAC9
Synonyms: HD7; HD7b; HD9; HDAC; HDAC7; HDAC7B; HDAC9B; HDAC9FL; HDRP; MITR
Vector: pCMV6-Entry (PS100001)
Fully Sequenced ORF: >SC331503 representing NM_001204144.
 Blue=Insert sequence Red=Cloning site Green=Tag(s)

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ATGATGAGCTCACCTGCACAGCCTGACCTCATGTGGAACCTGTACCATGGGTGCTATTCTGTGGCTGC
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AGTGCTTGTGTGGATGACACACTGGGACAAGTTGGGGCTGTGAAGGTCAAGGAGGAACAGTGGACAGT
GATGAAGATGCTCAGATCCAGGAAATGGAATCTGGGGAGCAGGCTGCTTTTATGCAACAGGTAATAGGC
AAAGATTTAGCTCCAGGATTTGTAATTAAGTCATTATCTGA
  
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Restriction Sites: SgfI-MluI



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ACCN:	NM_001204144
Insert Size:	1767 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:	NM_001204144.1
RefSeq Size:	4463 bp
RefSeq ORF:	1767 bp
Locus ID:	9734
UniProt ID:	Q9UKV0
Cytogenetics:	7p21.1
Protein Families:	Druggable Genome, Transcription Factors
MW:	65.5 kDa
Gene Summary:	<p>Histones play a critical role in transcriptional regulation, cell cycle progression, and developmental events. Histone acetylation/deacetylation alters chromosome structure and affects transcription factor access to DNA. The protein encoded by this gene has sequence homology to members of the histone deacetylase family. This gene is orthologous to the Xenopus and mouse MITR genes. The MITR protein lacks the histone deacetylase catalytic domain. It represses MEF2 activity through recruitment of multicomponent corepressor complexes that include CtBP and HDACs. This encoded protein may play a role in hematopoiesis. Multiple alternatively spliced transcripts have been described for this gene but the full-length nature of some of them has not been determined. [provided by RefSeq, Jul 2008]</p> <p>Transcript Variant: This variant (6) differs in the 5' UTR and coding sequence, the 3' UTR and coding sequence, and lacks an alternate in-frame exon compared to variant 1. The resulting isoform (6) is longer at the N-terminus, has a shorter and distinct C-terminus, and lacks an internal segment compared to isoform 1.</p>