

Product datasheet for **MC201670**

Suv39h1 (NM_011514) Mouse Untagged Clone

Product data:

| | |
|---------------------------|---|
| Product Type: | Expression Plasmids |
| Product Name: | Suv39h1 (NM_011514) Mouse Untagged Clone |
| Tag: | Tag Free |
| Symbol: | Suv39h1 |
| Synonyms: | AI852103; AL022883; DXHXS7466e; H3-K9-HMTase 1; KMT1A; mIS6 |
| Mammalian Cell Selection: | Neomycin |
| Vector: | PCMV6-Kan/Neo (PCMV6KN) |
| E. coli Selection: | Kanamycin (25 ug/mL) |



[View online »](#)

Fully Sequenced ORF: >BC023860 sequence for NM_011514
 CCACGCGTCCGCGCCCGCGCGCTCTCTTCTCGCGAGGCCGGTTAGGCCGAATGTTGTTAGCTGTGGA
 GAAAGATGGCGGAAAAATTTAAAAGGTTGCAGTGTGTGCTGTAATCTTCTTGGAAATCAACTGCAGGACCT
 GTGCCGACTAGCCAAGCTTTCTTGTCTGCCCTTGGTGTCTAAGAAGAATCTGTATGACTTTGAAGTT
 GAATACCTGTGTGATTATAAGAAGATCCGTGAGCAGGAGTATTACCTGGTTAAGTGGCGTGGGTATCCCG
 ACTCAGAAAACACCTGGGAGCCACGGCAGAATCTAAAATGTATACGAGTCTTAAGCAGTTCACAAGGA
 CTTAGAAAAGAGACTTGTCCGACGACACCGCCGGTCAAAGCCACCCAGGCATCTGGACCCAAACCTAGCC
 AATTACCTGGTGCAGAAGGCCAAGCAGAGCGGGCACTTCAGCGTTGGGAACAAGAGCTCAATGCCAAGC
 GCAGCCACCTGGGGCGGATCACCGTGGAGAATGAGGTAGACCTGGATGGCCCTCCAAGTCTTTGTCTA
 TATCAATGAGTATCGAGTTGGTGGGGCATCACCTCAACCAGGTAGCTGTTGGCTGTGAGTGCCAGGAC
 TGTCTGTTGGCACCCTGGAGGCTGTTGCCCTGGAGCATCCCTGCACAAGTTGCCTACAATGACCAAG
 GCCAGGTGCGACTGAAAGCTGGGAGCCACTACGAGTGCAACTCCCGCTGTTGCTGTGGCTATGACTG
 CCCAAACCGTGTAGTCCAGAAAGGCATCCGCTACGATCTCTGCATCTCCGCACTAATGATGGCCGAGGC
 TGGGGTGTCCGACGCTGAAAAGATCCGAAAAATAGCTTTGTTATGGAGTATGTGGGAGAGATTATA
 CCTCAGAGGAGGCAGAGCGGAGGGCCAGATCTACGACCGCCAGGGCCACCTACCTCTTTGACCTGGA
 CTACGTGGAAGACGTATATACCTGGATGCCGCTTATTATGGCAACATCTCTCATTTTGTCAACCATAGT
 TGTGATCCCAACCTGCAGGTGTACAACGTATTCATAGACAACCTTGATGAGCGACTACCCCGCATCGCAT
 TCTTTGCCACAAGAACCATCTGGGCGGGCGAGGAGCTCACCTTTGATTACAACATGCAAGTGGACCCCGT
 GGACATGGAGAGTACCCGAATGGACTCCAACCTTTGGCCTGGCTGGGCTCCCCGGCTCCCCAAGAAACGA
 GTCCGATTGAATGCAATGTGGGACAACGGCTTGCCGAAAAACCTCTTCTAGCCCTGAGAAGTCTGAG
 GCCAGACTAACTGAAGGGGCTGAAGCCACCTTCTCTCTACAGCTACCTCTTGTCAAGGATGACCAT
 CAATCAGAGCCTTGTCTGCCTCCACTTGTCTCACCTACCCTAACCTGCTTAGGGTCAGGGCTGTTGTG
 AGGACTAACTCCGGGTACCCTTTTCTGTTCTTTCCCTGTTCCAGGCCCATCAGGCATTGCATTAA
 AACTCCCAGCCCCATTTTCAGAAACATATTTTTCACATCATGATTCCTAGAGTTGGAATTCATGTCATA
 TAATGGAGGTCCAGATTGAGGAACCTCGGCTGTAAAACAGATTCTTTGTTTTGGACAGCATCTCTGCAGCT
 CTATGTAGTAAGTCTGGTGTGGACCGTTAATCTTCTGCTCAGCCTTCTCATGATGAGATTGTAGG
 TGTAAACCCAGCTAAGATTTTTGTTCTAATTGCATCTGCTTCTGCTTGGAGCTTGTGTGACCTGTTGC
 AGGCCTCTTCACTACTAATGGTATGAGGAAGCAACCCCTGGCAGACAGACATCAGAGCTGAGATACC
 AGCCTAACATCAAGCTGGATCAGCAACCCAGAGCCTTGTACTCAGGAAAGAAAAGGCAATCTTCAGAG
 CTGGGAGATAAGGCTGGTCCGCTCTTTGTGCTTTTGATGCTGGCTGGTATTAACCTAAGACCTATAGG
 GTCTCAACAGTTGCAAGTCTGAAAAGTAGTTGCCCAAATGCCATCAGAATGGGGATGGAGTAAATACCTC
 TTTGAAAGCCCCACAGAAAGTTAGAAGTAAAGTTTACCATCAGGAAGTACAGTGCCTGGACTTGCTGGAA
 ACCCAGCCTTGGCATTGATGGCCACTAGAAGCTAGAAAGCTGGAACCAAGATCTAGGTATTTCTTAGA
 TAGCACTTAAGACAGTAATGTGCATCGACTAGAAGGCATGTGATGCCAGGCACTTGGTAGAGCACCTGG
 TCCATACAGATTGTCTCAGGGAAGCCTTGAAAACCACAAAGGTGGAGCCAGAAAAAGCCCATGTGACA
 GAAGGCAATGTCTAGGCCAAAAATACTTGTGAGTCCAAGTATTCACCTGGGTCCACTTGTCTCAGTTAA
 CTGCCTAGAAATGTACAAAAGGCAAGATTCTGATGGCTGCCTTGCCCTGCTTCCCACCTCCAGGAA
 GCCTTTCTGACTTCTGTGCCAGAGTGCCCTATGTGAACTCTGTACCCTGCTACCAGATGCCAGGTC
 TGTGTGTATTTGTATATATGTTTCTGCCATACTTCCCAGGCTGACCTTCCAGGCATGGACTGAATC
 TGGTCTCTCTGTACCCCTCAGCCCTCCCTAGCCTGGAGTGCACACCAATAAACTGTGTTGTTGAGTTAA AAAAAAAAAAAAA

Restriction Sites: RsrII-NotI
ACCN: NM_011514
Insert Size: 1239 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: | BC023860 , AAH23860 |
| RefSeq Size: | 2743 bp |
| RefSeq ORF: | 1239 bp |
| Locus ID: | 20937 |
| UniProt ID: | O54864 |
| Cytogenetics: | X 3.64 cM |
| Gene Summary: | <p>Histone methyltransferase that specifically trimethylates 'Lys-9' of histone H3 using monomethylated H3 'Lys-9' as substrate. H3 'Lys-9' trimethylation represents a specific tag for epigenetic transcriptional repression by recruiting HP1 (CBX1, CBX3 and/or CBX5) proteins to methylated histones. Mainly functions in heterochromatin regions, thereby playing a central role in the establishment of constitutive heterochromatin at pericentric and telomere regions. H3 'Lys-9' trimethylation is also required to direct DNA methylation at pericentric repeats. SUV39H1 is targeted to histone H3 via its interaction with RB1 and is involved in many processes, such as repression of MYOD1-stimulated differentiation, regulation of the control switch for exiting the cell cycle and entering differentiation, repression by the PML-RARA fusion protein, BMP-induced repression, repression of switch recombination to IgA and regulation of telomere length. Component of the eNoSC (energy-dependent nucleolar silencing) complex, a complex that mediates silencing of rDNA in response to intracellular energy status and acts by recruiting histone-modifying enzymes. The eNoSC complex is able to sense the energy status of cell: upon glucose starvation, elevation of NAD(+)/NADP(+) ratio activates SIRT1, leading to histone H3 deacetylation followed by dimethylation of H3 at 'Lys-9' (H3K9me2) by SUV39H1 and the formation of silent chromatin in the rDNA locus. Recruited by the PER complex to the E-box elements of the circadian target genes such as PER2 itself or PER1, contributes to the conversion of local chromatin to a heterochromatin-like repressive state through H3 'Lys-9' trimethylation.[UniProtKB/Swiss-Prot Function]</p> |