

Product datasheet for **SC303814**

EHMT2/G9A (EHMT2) (NM_006709) Human Untagged Clone

Product data:

Product Type: Expression Plasmids
Product Name: EHMT2/G9A (EHMT2) (NM_006709) Human Untagged Clone
Tag: Tag Free
Symbol: EHMT2
Synonyms: BAT8; C6orf30; G9A; GAT8; KMT1C; NG36
Mammalian Cell Selection: None
Vector: [pCMV6-XL6](#)
E. coli Selection: Ampicillin (100 ug/mL)

Fully Sequenced ORF: >OriGene sequence for NM_006709 edited
 GCGCAAGCGGCGATGGCGGCGGGGAGCTGCAGCGGCGGCGCCGAGGGGGAG
 GCCCCCGCTGAGATGGGGGCGCTGCTGCTGGAGAAGAAACCAGAGGACCCAGAGAGA
 GTTCATGGCTCTTTGGGGGACACCCTCGTAGTGAAGAAACCCTGCCAAGGCCACCCCC
 GACTCCCTGGAGCCTGCTGGCCCTCATCTCCAGCCTCTGTCACTGTCACTGTTGGTGAT
 GAGGGGGCTGACACCCTGTAGGGGCTACACCACTCATTGGGGATGAATCTGAGAATCTT
 GAGGGAGATGGGGACCTCCGTGGGGGCCGATCCTGCTGGGCCATGCCACAAAGTCATTC
 CCCTCTTCCCCCAGCAAGGGGGTTCTGTCTAGCCGGGCAAGATGTCAATGACAGGG
 GCGGGAAAATCACCTCCATCTGTCCAGAGTTGGCTATGAGGCTACTGAGTATGCCAGGA
 GCCCAGGGAGCTGCAGCAGCAGGGTCTGAACCCCTCCAGCCACCAGAGCCAGAGGGA
 CAGCCCAAGGTCCACCGAGCCCGCAAACCATGTCCAACCCAGGAAATGGACAGCCCCCG
 GTCCCTGAGAAGCGGCCCTGAAATACAGCATTTCGCGATGAGTGATGATGTCCACTCA
 CTGGGAAAGGTGACCTCAGATCTGGCCAAAAGGAGGAAGCTGAACTCAGGAGGTGGCCTG
 TCGGAGGAGTTAGGTTCTGCCGGCGTTCAGGAGAAGTGACCCTGACGAAAGGGGACCCC
 GGGTCCCTGGAGGAGTGGGAGACGGTGGTGGTGATGACTTCAGTCTCTACTATGATTCC
 TACTCTGTGGATGAGCGCGTGGACTCCGACAGCAAGTCTGAAGTTGAAGCTCTAACTGAA
 CAACTAAGTGAAGAGGAGGAGGAGGAAGAGGAGGAAGAAGAAGAAGAGGAAGAGGAGGAG
 GAAGAGGAAGAAGAAGAGGAAGATGAGGAGTCAAGGAATCAGTCAGATAGGAGTGGTTCC
 AGTGGCCGCGCAAGGCCAAGAAGAAATGGCGAAAAGACAGCCCATGGGTGAAGCCGTCT
 CGGAAACGGCGCAAGCGGGAGCCTCCGCGGGCCAAGGAGCCACGAGGAGTGAATGGTGTG
 GGCTCCTCAGGCCCAAGTGAGTACATGGAGTCCCTCTGGGGTCCCTGGAGCTGCCACG
 GAGGGGACCTCTCCCCAACCCAGCTGGGGTGTCCAATGACACATCTTCGCTGGAGACA
 GAGCGAGGGTTTGGAGGTTGCCCTGTGCAGCTGCCGATGGAGGCACCCAAGATTGAC
 CGCATCAGCGAGAGGGCGGGGCACAAGTGCATGGCCACTGAGAGTGTGGACGGAGAGCTG
 TCAGGCTGCAATGCCGCATCCTCAAGCGGGAGACCATGAGGCCATCCAGCCGTGTGGCC
 CTGATGGTGTCTGTGAGACCCACCGCGCCCGCATGGTCAAACACCACTGCTGCCGGGC
 TGGCGTACTTCTGCACGGCGGCACCTTCTGGAGTGCCACCCTGACTCCGTGTGGCC



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CACCGCTTCCACAAGGCCTGTGTGTCTCAGCTGAATGGGATGGTCTTCTGTCCCCACTGT
GGGGAGGATGCTTCTGAAGCTCAAGAGGTGACCATCCCCGGGGTGACGGGGTGACCCCA
CCGGCCGGCACTGCAGCTCCTGCACCCCAACCCCTGTCCCAGGATGTCCCAGGAGAGCA
GACTTCTCAGCCAGTGCCCGGATGCGAGGGCATGGGAACCCCGCGCCCGCCCTGC
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TGCCCTTTCAGCCGTGGGGCTGCCACTGGGGCCAGGCCGGGAGGCCCTGGAAAAGGCCCTG
GTCATCCAGGAGTCAGAGAGGCGGAAGAAGCTCCGTTTCCACCCTCGGCAGTTGTACCTG
TCCGTGAAGCAGGGCGAGCTGCAGAAGGTGATCCTGATGCTGTTGGACAACCTGGACCCC
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TCCGTGGAGATCTGCCATGTGCTGCTGCAGGCTGGAGCCAACATAAATGCAGTGGACAAA
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ATGGTGCAGCGTGGTGGCTGTGTCTATAGCAAGGAGGAGGACGGTTCCACCTGCCTCCAC
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GTGTGGTTTGGCGTTCAACTCAACCGCAAGCTCCGACTTGGGGTGGGAAATCGGGCCATC
CGCACAGAGAAGATCATCTGCCGGGACGTGGCTCGGGCTATGAGAACGTGCCATTCCC
TGTGTCAACGGTGTGGATGGGGAGCCCTGCCCTGAGGATTACAAGTACATCTCAGAGAAC
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GACAAGGATGGGCGATTGCTCCAGGAATTAACAAGATTGAGCCTCCGCTGATTTTCGAG
TGTAACCAGGCGTGTCTATGCTGGAGAACTGCAAGAACCGGGTGCTACAGAGTGGCATC
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ACCATCCACAGGGGACCTTCTGCGAGTATGTCGGGGAGCTGATCTCTGATGCTGAG
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TACTGCATAGATGCCGTTACTATGGCAACATCAGCCGTTTCATCAACCACCTGTGTGAC
CCCAACATCATTCCCGTCCGGTCTTTCATGCTGCACCAAGACCTGCGATTTCCACGCATC
GCCTTCTCAGTTCGAGACATCCGGACTGGGGAGGAGCTAGGGTTTGACTATGGCGAC
CGTCTTGGGACATCAAAGCAAATATTTACCTGCCAATGTGGCTCTGAGAAGTGAAG
CACTCAGCCGAAGCCATTGCCCTGGAGCAGAGCCGTCTGGCCCGCTGGACCCACACCTT
GAGCTGTGCCGAGCTCGGCTCCCTGCCCCCTGTCAACACATGAGAACGGACCACACC
TCTCTCCCAGCATGGATGGCCACAGCTCAGCCGCTCCTCTGCCACCAGCTGCTCGCAG
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GAGATCCCAGCCAGGCCCTGGAGGTCTGACAGCCCTCCCTCCCAGAGCTGGTTCCCTCCC
TGGGAGGGCAACTTCAGGGCTGGCCACCCCGTGTTCCCATCCTCAGTTGAAGTTTGA
TGAATTGAAGTCGGGCTCTATGCCAACTGGTTCCTTTTGTCTCAATAAATGTTGGGTT
TGGTAAAAAAAAAAAAAAAAA

5' Read Nucleotide Sequence: >OriGene 5' read for NM_006709 unedited
 ACCCGCCCCGTCTGAGCAACTGGGCGGTAGGCGCTGCACTGTGTTGGGAGGTTCTATTAT
 AAGCAGAGCTCATTTAGGTTGACTATAGAATACAAGCTACTCTGTTCTTTTTGCAGCG
 GCCGCGAATTTCCGGCAGGAGGGCGCAAGCGCGATGGCGGCGGCGGGAGCCGAGCG
 GCGGCGGCGCCGAGGGGGAGGCCCCCGCTGAGATGGGGGCGCTGCTGCTGGAGAAGGAA
 ACCAGAGGAGCCACCGAGAGAGTTCATGGCTCTTTGGGGACACCCCTCGTAGTGAAGAA
 ACCCTGCCCAAGGCCACCCCGACTCCCTGGAGCCTGCTGGCCCTCATCTCCAGCCTCT
 GTCAGTGTCACTGTTGGTATGAGGGGGCTGACACCCCTGTAGGGGCTACACCACCTATT
 GGGGATGAATCTGAGAATCTTGAGGGAGATGGGACCTCCGTGGGGGCGGATCCTGCTG
 GGCCATGCCACAAAGTCATTCCTCTTCCCCAGCAAGGGGGTTCTGTCTAGCCGG
 GCCAAGATGTCAATGACAGGGGCGGAAAATCACCTCCATCTGTCCAGAGTTTGGCTATG
 AGGCTACTGAGTATGCCAGGAGCCAGGGAGCTGCAGCAGCAGGGTCTGAACCCCTCCA
 GCCACCAGAGCCAGAGGGACAGCCCAAGGTCCACCGAGCCGAAAACCATGTCCAAA
 CCAGGAAATGGACAGCCCGGTCCCTGAGAAGCGGCCCTGAAATACAGCATTTCGC
 ATGAGTGTATGATGCCACTCACTGGGAAAGGTGACCTCAGATCTGCCAAAAGAGAAGCT
 GAACTCAGAGTGGCTGTGAGAGTAGTCTGCCGCGTTACAGAGAGTGACCCTGACAAA
 GACCCGGTCTGAGATGGAGACGTGTGGGGTGTGACTCAGTCTCTACTATGATCCTACT
 CTGTGGATGAGCCGTGGACTCGACG

3' Read Nucleotide Sequence: >Forward primer walk for NM_006709 unedited
 TATTGCAGATGTTGTTGGAGACCTGCTCTAGCTCCACTAGCCATGTTGCGCCAGCTTCA
 GCATCCGGTGTGGTATTGACAAGGATGGGCGATTGCTCCAGGAATTTAACAAGATTGAG
 CCTCCGCTGATTTTCGAGCGTAACCAGGCGCTCATGCTGGAGAAACTGCAAGAACCGG
 GTCGTACAGAGTGGCATCAAGGTGCGGCTACAGCTCTACCGAACAGCCAAGATGGGCTGG
 GGGTCCGCGCCTGCAGACCATCCCACAGGGACCTTCATCTGCGAGTATGTCGGGGAG
 CTGACCTCTGATGCTGAGGCTGATGTGAGAGAGGATGATTCTTACCTCTCGACTTAGAC
 AACAAGGATGGAGAGGTGACTGCATAGATGCCGTTACTATGGCAACATCAGCCGCTTC
 ATCAACCACCTGTGTACCCCAACATCATTCCCGTCCGGGTCTTCATGCTGCACCAAGAC
 CTGCGATTTCCACGCATCGCCTTCTTCAAGTCCCGAGACATCCGACTGGGGAGGAGCTA
 GGGTTTGACTATGGCGACCGCTTCTGGGACATCAAAGCAAATATTTACCTGCCAATGT
 GGCTCTGAGAAGTGAAGCACTCAGCCGAAGCCATTGCCCTGGAGCAGAGCCGTCTGGCC
 CGCCTGGACCCACACCCTGAGCTGCTGCCCGAGCTCGGCTCCCTGCCCTGTCAACACA
 TGAGAACGGACCACACCCTCTCTCCCAGCATGGATGGCCACAGCTCAGCCGCTCCTCT
 GCCACCAGCTGCTCGCAGCCATGCCTGGGGTGTGCCATCTCTCTCCCACCACCCTT
 TCACACATTCCTGACCAGAGATCCAGCCAGGCCCTGAGTCTGACAGCCCTCCCTCCCA
 GAGCTGTTCCCTGGGAGGGCAACTT

Restriction Sites: Please inquire
ACCN: NM_006709
Insert Size: 4000 bp

OTI Disclaimer: 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 custsupport@origene.com or by calling 301.340.3188 option 3 for pricing and delivery.

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. [More info](#)

OTI Annotation: The open reading frame of this TrueClone was fully sequenced and found to differ from the protein associated to this reference by a single amino acid.

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:

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_006709.2](#), [NP_006700.2](#)

RefSeq Size: 3994 bp

RefSeq ORF: 3633 bp

Locus ID: 10919

UniProt ID: [Q96KQ7](#)

Cytogenetics: 6p21.33

Domains: SET, ANK, PreSET, Pre-SET

Protein Families: Druggable Genome

Protein Pathways: Lysine degradation

Gene Summary:

This gene encodes a methyltransferase that methylates lysine residues of histone H3. Methylation of H3 at lysine 9 by this protein results in recruitment of additional epigenetic regulators and repression of transcription. This gene was initially thought to be two different genes, NG36 and G9a, adjacent to each other in the HLA locus. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Jan 2016]

Transcript Variant: This variant (2, also known as NG36/G9a) uses an alternate 5' UTR and 5' coding region, contains an alternate in-frame exon in the 5' coding region, and lacks an alternate in-frame exon compared to variant 3. It encodes isoform a, which is shorter and has a distinct N-terminus, compared to isoform c.