

## Product datasheet for MR223799

### Kmt2e (NM\_026984) Mouse Tagged ORF Clone

#### Product data:

Product Type:	Expression Plasmids
Product Name:	Kmt2e (NM_026984) Mouse Tagged ORF Clone
Tag:	Myc-DDK
Symbol:	Kmt2e
Synonyms:	1810033J14Rik; 9530077A04Rik; D230038D11Rik; MII5; NKp44L
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
ORF Nucleotide Sequence:	>MR223799 representing NM_026984, <b>codon optimized</b> . Due to the complexity of NM_026984, the ORF clone is codon optimized for mammalian Expression. The nucleotide sequence differs from the reference sequence, yet the amino acid sequence remains identical.

Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC  
GCC**CGGATCGCC**

ATGAGCATAGCGATCCCATTGGGGTTGATACAACAGAAACATCCTACTTGGAAATGGCTGCAGGCTCAG  
AACCGGAATCCGTAGAAGCTAGCCCTGTGGTAGTTGAGAAATCCAACAGTTTTCCCACCAAGTTATACAC  
CAGCAGCTCCCACCACTCACACAGCTACATCGGTCTGCCCTATGCGGACCATAATTATGGTGCTCGTCTCT  
CCTCCAACACCTCCAGCATCCCCCTCCATCAGGTCTCATTAGCAAAAACGAAGTAGGCATATTTACCA  
CTCCTAATTTTGATGAACTTCCAGTGCTACTACAATCAGCACATCTGAAGATGGAAGTTACGGTACTGA  
TGTAACCAGGTGCATATGTGGTTTTACACATGATGACGGATACATGATCTGTTGTGACAAATGCAGTGTT  
TGGCAACATATTGACTGCATGGGATTGACAGGCAGCATATTCCTGATACATACCTATGTGAACGTTGTC  
AGCCCAGGAGTTTGATAAAGAGAGGGCAGTGCTACTGCAACCGCGAAAAGAGAAAATATGTCAGATGG  
TGATACCAGTGCAACTGAAAGTGGTGATGAAGTTCCTGTGGAATTATATACTGCATTTCCAGCATACTCCA  
ACTTCAATCACTTAACTGCTTCCAGAGTTCCTCAAGGTTACTGATAAAAAGAGGAAAAAAGTGGGGAGA  
AAGAACAAAACTTTTCAAATGTAAAAAGCCTTTCGTGAAGGATCTAGGAAATCATCAAGGGTTAAGGG  
TTCAGCACCAGAAATTGATCCTTCATCTGATAGTTCAAATTTTGGTGGGAAACAAAATCAAAGCATGG  
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CTCAAAGACTCGGTAGTGGGAATGACAGCAAAGACATGAATAAATCAGAATTGAGTACCAACAACACTCACT  
CTTCAGACCTCCTGTAGAGAGCCATATACAGAAGAATAAAAAAATCTGAAATCTGCCAAAGACTTGCCCT  
CCTGATGCACTTATCATTGAATATAGAGGGAAGTTTATGCTGAGAGAACAGTTTGAAGCAAATGGATATT



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TCTTTAAAAGGCCATACCCTTTTGTATTCTACTCTAAGTTCCATGGGCTTGAATGTGTGTTGATGC  
 GAGGACTTTTGGGAATGAGGCTCGATTCACTCAGGCTTCTGTACACCGAATGCAGAGGTGAGGCATGAA  
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 GTCAAACAACCAGGAACCAGATTTTATTGATGATATGGAAGAAAAAAGCTCTATTAGCAATGAAGTGGAA  
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 AGCTAAGACAGAAGTTAAGCCTGAATGCAAGAGTTCGAGGTCATCGCCGATGCCGAAGTGGTGCAGGAA  
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 TTTCTCGTAGTAGGACTCACATTGGACAGCAGCGTCGGAGACACAGAACTGTCAGCATGTGTTTCCAGCAT  
 ACCGCCATCTTCTCTGACATCGAAGTTCTGTCAACAGAACGAGATTGAAAATACTGTACTTGTCTATA  
 GAACCAGAACTGAACTGCAGTAGCAGAAATAATTCTGAAGCGGAAGTCCAGCACTTAATAAGTGTG  
 CCACAAAGTACCCAAAACAAGAGCACTTTGGTCAATGAATGGTTAAGTGAGAAGAATGAGAAGACAGG  
 AAAACCTTCAGACAGCCTTTAGAAAAGGCTCTGCGCATAACTACAGATCCAGAGGTGTTAGCTACACAG  
 CTCAATTTCTTTGCCAGGTCTCACTTACAGCCCCATGTATACTCTACTCCTAAGCATTATATTAGATTTA  
 CATCACCATTCTTTTCGAAAAAAGAGAAAGAAAAGAACTACTGAAAACATTTCTGGCTCATGCAAAAA  
 GCGGTGGCTGAAGCAAGCCCTGGAGGAAGAAAATCAACAATTTTACATAGATATCATTACCTTGTGAG  
 GAAAGATCCAGGAGTCTACAGTTAATGGTGAAAACAAAAGTCCACTTCTGCTAAGCGACAGCTGCTCCC  
 TCCCAGATTTAACACACCACTAAAAAGCGAAGACTTTACCAATTGTTAGACTGCTTACTCAGAAAAG  
 CTCCACTCTACCCCTTACCATATGCTACACCAACTCACACAGACATCACTCCTACGACCCAGCACTTT  
 GCCACTCCCCACGGATCAAATCAGATGATGAAACTTACAGAAATGGTTATAAACCCATATATTCACCAG  
 TGACCCCTGTAACCCCTGGCACACCAGGAAACACCATGCACTTTGAGAATATTTCTTCCCCAGAAAGTTC  
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 CCTTTTAGAAATTCAAATTAAGTGAAGTGGGCTGCAAGAAAATAAGACTATTGGTTATACCAGCCCTA  
 GGAGCAGGACTGAGGTTAACAGGCTTGGCCAGGGGAGAGGAGTCTGTGTGAGACCTTCAACTGGGACT  
 CGATGCAGTCGAGCCAGTGCCTTACAGAAATCCATGGAAACACCAGCCCATGACAGGACTGAGCCAGC  
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 CAGGAGTTAACTTCTCAGTAACTCCAAGTGGAGGACTTAACACCTCTCATCAGTTGGAGACTGGAGG  
 CGGCTTCCGAGTAAGTGAGTCCAAGTGCCTGATTCAGCAGGATGACACTAGAGGCATGTTTCTGGGAGCG  
 GCTGTCTTTTGTACTTCTGAAGACGGGCTTGCCTCTGGTTTTGGACGGACTGTTAATGACATTTGATCG  
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 ACTCACAAAACCAGATAGCCACTGGGAGGCAACTGCCACTGTATCAGAAGCTGACAACAGTGTTCACCAA  
 AACCCAGAGCCTCAGCACAGACAGCTGTCAAGTAAACCCAGCAGCTTCTCAGAACCATGCTCCTCAGG  
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 TCCGTACCTCACACAGAAAACCTCCCAAGTCTCCACACTCACACCCCGTGCAGCATGTTTACCTC  
 TCGCCAAAGCCTCCTTACAGCACTTAGGATCTCCCTTACAGCCTCACCATTACAATCACCTCAAGTTG  
 GAACTCCTCAACGAGAGACTCAGCGCAACTTTTACGCCGACGCCAGAATCTGCAGGCAACCCCAAGCA  
 GGCTACCTCAGGAGCTCTTTTACTCAAACCTTTCAGGGCAGTCCCGCAACTTATCCAGTTCAAT  
 CAGCAAAGCCTCAATAGTACGGCCCCGCTCCACTCCACCGCCGCTCTAGTAGTTATTATCAAAAACC  
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AAGCGGGCCTAATCAAGCCCTGCCTGGCTCTACTTCACAGCAGTCCGTGCCTGGGCACCACGTACACCT  
 GGGCACTTCTGCCATCCCAGAACCCACAATACACCATCAACCTGCCGCGGCCGCCGTCTGCCACCTC  
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 CCCACCCCGCCGCTAGCGTGTGGTTCAGCGGCATCACTCCGCCAGTGGTCAGGCCCTGCACCACCT  
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 CCCACCACACCACCTGGGCCCGGCCCTCAACACCAGCCATCTGGCACTGGCCCTCACTGTCTCTCCC  
 CGTCGAGGGCCACACTTGCAGCCTCAGGGGCCAAATAGCATACCTACTCTACAGCTTCAGTTTTTGT  
 CCTCACCCACCCAGGTAGCGTGGCGCTGCCACACGGCGTGCAGGGTCTCAGCAGGCCTCCCCTGTGC  
 CCGCGCAGATTCCAATCCACAGGCCCAAGTGCCTCCCACTTTCCAGAATAACTATCACGGTTCCGGATG  
 GCAC

ACGCGTACGCGGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCTGGATT  
 ACAAGGATGACGACGATAAGGTTTAA

**Protein Sequence:**

>MR223799 representing NM\_026984  
 Red=Cloning site Green=Tags(s)

MSIAIPLGVDTTETSYLEMAAGSEPESEVESPVVVEKSNFPHQLYSSSHSHSYIGLPYADHNYGARP  
 PPTPPASPPPSGLISKNEVGIFFTPNFDETSATTISTSEDSYGTVDVTRICIGFTHDDGYMICCDKCSV  
 WQHIDCMGIDRQHIPDTYLCERCQPRSLDKERAVLLQRRKRENMSDGDTSATESGDEVPVELYAFQHTP  
 TSITLTASRVPKVTDKRRKKSQEKEQNFQSKCKKAFREGSRKSSRVKGSAPIDPSSDSSNFVWETKIKAW  
 MDRYEEANNNQYSEGVRQEAQRLAQRLLGSGNDSKDMNKSELSTNNSLFRPPVESHQKKNKILKSAKDL  
 PDALIEYRGKFLRQFEANGYFFKRPYFVLFYSKFHGLEMCVDARTFGNEARFIRRSCTPNAEVRHE  
 IEEGTIHLIYISIQSIPKGEITIAFDYGNCKYKVDCAKLKENPECPVLRKRSSESTENINSGYETRRK  
 KGKKEKDTSKEDIQNQNTLDCEGTNNKIRSPETKQRKLSPLRLSVSNNQEPDFIDDMEEKTPIISNEVE  
 MESEEQIAERKRKMTREERKMEAILQAFARLEKREKRREQALERISTAKTEVKPECKESQVIADAEVVQE  
 QKKEETAIKPAAAKVNRKQKRSFSRSRTHIGQRRRHRTVSMCSDIPSSPDIEVLQQNEIENTVLAIE  
 EPETETAIAEIIPEAEVPALNKCPKYPKTKKHLVNEWLSEKNEKTGKPSDSLSEPLRITTDPEVLATQ  
 LNSLPGLTYPHYSTPKHYIRFTSPFLSEKRRKETTENISGCKKRWLKQALEENSTILHRYHSPCQ  
 ERSRSPVNGENKSPLLLSDSCSLPDLTTPKRRRLYQLLDTAYSESSTPTSPYATPTHTDITPTDPAF  
 ATPPRIKSDDETYRNGYKPIYSPVTPVTPGTPGNTMHFENISSPESSPEIKRCTYNQEGYDRPSNMLTG  
 PFRNSNLTEGLQEIKTIGYTSRPRSRTEVNRPCPGEKESVSDLQLGLDAVEPAALQKSMETPAHDRTEPS  
 NQLDSTHSGRGTMYSSWVKSPDRGTGVNFSVNSNLRDLTPSHQLETGGFRVSESKLIQQDDTRGMFLGA  
 AVFCTSEDGLASGFGRVNDNLIDGCTPQNPQKVVSLLEYRKRQREARKSGSKPENFALISVSPHPS  
 GSLSSSGDGCVHSENGEQAENQASLPLPPAAAAAATAAAAYSASSEEGSSNCPVKDANSSEKKDPEVQ  
 WTAQSVQVRSYQRAALLSDHRKDKDSGGESPCVSCSPSHVQSPSSSHNHIPQVHAQSLAPSLSEL  
 MADPDAEGTEATSTSECPSPDTSQSPSKTSKPGSPGPIPAQSHGKILTKPDSHWATATVSEADNSVHQ  
 NPEPQHRQLSSNTPALSQNHAPQAHALSANDQLPQKLPSAPTCLHCPPSPHTENPPKSSTPHTPVQHGYL  
 SPKPPSQHLGSPFRPHHSQSPQVGTQRETQRNFYAAAQNLQANPQQATSGALFTQTPSGQSSATYSQFN  
 QQSLNSTAPPPPPPPSSYYQNPQSANFQNYNQLKGSLSQQTVFTSGPNQALPGSTSQQSVPGHHVTP  
 GHFLPSQNPТИHQPAAAAAVPPPPPPAPGPHLIQQPSSHQQHVAHVGVGPHAVTPGSHIHSQTAGH  
 HLPPPPPPGPAPHHPPPHPTGLQSLQAQHQHVNSAPPPPPPPPPASVLVSGHHSASGQALHHP  
 PHQGPPLFPASAHPAVPPYPSQATHHTLGGPQHQPSTGPHCLPVAGPHLQPGQNSIPTPTASGFC  
 PHPHGVSVALPHGVQGPQASPVPAQIPIHRAQVPPPTFQNNYHSGSWH

TRTRPLEQKLISEEDLAANDILDYKDDDDKV

**Restriction Sites:**

Sgfl-MluI

**Cloning Scheme:**


**ACCN:** NM\_026984

**ORF Size:** 5604 bp

**OTI Disclaimer:** 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:** This clone was engineered to express the complete ORF with an expression tag. Expression varies depending on the nature of the gene.

**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\\_026984.1](#), [NP\\_081260.1](#)

**RefSeq Size:** 7229 bp

**RefSeq ORF:** 5607 bp

**Locus ID:** 69188

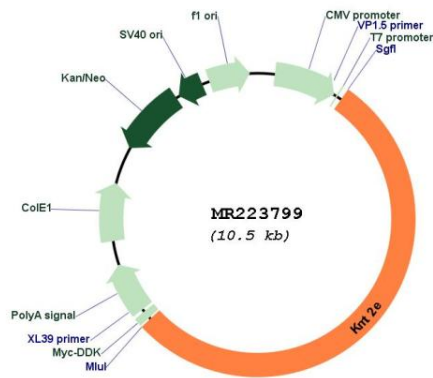
UniProt ID: [Q3UG20](#)

Cytogenetics: 5 A3

MW: 204.5 kDa

**Gene Summary:** Associates with chromatin regions downstream of transcriptional start sites of active genes and thus regulates gene transcription (By similarity). Chromatin interaction is mediated via the binding to tri-methylated histone H3 at 'Lys-4' (H3K4me3) (By similarity). Key regulator of hematopoiesis involved in terminal myeloid differentiation and in the regulation of hematopoietic stem cell (HSCs) self-renewal by a mechanism that involves DNA methylation (PubMed:18854576, PubMed:18952892, PubMed:18818388). Also acts as an important cell cycle regulator, participating in cell cycle regulatory network machinery at multiple cell cycle stages including G1/S transition, S phase progression and mitotic entry (PubMed:19264965). Recruited to E2F1 responsive promoters by HCFC1 where it stimulates tri-methylation of histone H3 at 'Lys-4' and transcriptional activation and thereby facilitates G1 to S phase transition (By similarity). During myoblast differentiation, required to suppress inappropriate expression of S-phase-promoting genes and maintain expression of determination genes in quiescent cells (PubMed:19264965).[UniProtKB/Swiss-Prot Function]

### Product images:



Circular map for MR223799