

MLL1 (D6G8N) Rabbit mAb (Carboxyterminal Antigen)



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| Applications: W, IP, IF-IC | Reactivity: H M R Mk | Sensitivity: Endogenous | MW (kDa): 180 | Source/Isotype: Rabbit IgG | UniProt ID: #Q03164 | Entrez-Gene Id: 4297 |
|-------------------------------|--------------------------------|--|---|---|--|---|
| Product Usage Information | 2 | Application Western Blotting Immunoprecipitation Immunofluorescence | (Immunocytochem | iistry) | | Dilution 1:1000 1:50 1:200 |
| Storage | | Supplied in 10 mM soc 0.02% sodium azide. S | dium HEPES (pH 7.5 tore at –20°C. <i>Do r</i> | 5), 150 mM NaCl, 100 μg/ not aliquot the antibody. | ml BSA, 50% glyce | rol and less than |
| Specificity/Ser | nsitivity | MLL1 (D6G8N) Rabbit mAb (Carboxy-terminal Antigen) recognizes endogenous levels of total MLL1-C protein. | | | | |
| Source / Purifi | cation | Monoclonal antibody i carboxy terminus of h | is produced by imn uman MLL1 proteii | nunizing animals with rea. n. | combinant protein | specific to the |
| Background | | The Set1 histone meth histone methyltransfe transcriptional co-activ Set1-related proteins: COMPASS-like complex found in distinct prote CXXC1 and DPY30, whi methyltransferase activ menin (6). | nyltransferase prote rase complex, whic vator (1). While yea SET1A, SET1B, MLL xes and methylate in complexes, all o ich are required foi ivity (2-6). MLL1 an | ein was first identified in th methylates histone H3 st contain only one know 1, MLL2, MLL3, and MLL- histone H3 at Lys4 (2,3). f which share the commo r proper complex assemi d MLL2 complexes conta | yeast as part of the at Lys4 and function of Set1 protein, mar at all of which assent These Set1-related on subunits WDR5, oly and modulation in the additional protection | e Set1/COMPASS ons as a mmals contain six mble into proteins are each RBBP5, ASH2L, of histone rotein subunit, |
| | | MLL1 functions as a m proper expression of H cleaved by the taspase (ML1-C) fragments, b N, MLL1-C, WDR5, RBE complex, which is recr transcriptional initiatio characterized and asso translocation partners of these partners are r AF4, AF9 and ENL are a ENL all interact with th normally regulated by transcriptional elonga translocation partners specific gene expression | aster regulator of Hox genes (7,8). ML a 1 threonine endo oth of which are su 3P5 and ASH2L defi uited to target gen on (11). At least 60 ociated with variou s include AF4, AF9, I nuclear proteins th all components of the histone H3 lysing promoter-proxima- tion occurring in re- s with SEC and DOT on programs by co | both embryogenesis and L1 is a large, approximal peptidase to form N-terr ibunits of the functional ne the core catalytic con les and methylates histo different MLL1 translocal s hematological maligna ENL, AF10, ELL and AF6 (3 at function to positively to the super elongation cor e 79 methyltransferase D al pausing, with the releat esponse to proper stimul 1L suggest that MLL1-fu nstitutively activating tra | I hematopoiesis, ar tely 4000 amino ac ninal (MLL1-N) and MLL1/COMPASS co uponent of the MLL ne H3 lysine 4 to re- tion partners have ncies. The most co 3,12,13). With the e regulate transcripti nplex (SEC), while A OT1L. Many MLL1 se of RNA polymer i (14). The associati sion proteins may f nscriptional elonga | nd is required for id, protein that is C-terminal MLL1 mplex (9,10). MLL1- .1/COMPASS gulate been molecularly mmon xception of AF6, all onal elongation. .F4, AF9, AF10 and target genes are ase and on of MLL1 function to sustain ation. |
| Background R | eferences | Miller, T. et al. (2001) Shilatifard, A. (2008) Tenney, K. and Shila Lee, J.H. and Skalnik Lee, J.H. et al. (2007) Hughes, C.M. et al. (Teissenberg, J.C. and Smith, E. et al. (2011) Takeda, S. et al. (2001) Yokoyama, A. et al. Dou, Y. et al. (2006) Yip, B.H. and So, C. Neff, T. and Armstr |) Proc Natl Acad Sc) Curr Opin Cell Bio tifard, A. (2005) J C c, D.G. (2005) J Biol) J Biol Chem 282, 1 (2004) Mol Cell 13, Shilatifard, A. (201) Genes Dev 25, 66)6) Genes Dev 20, 2 (2002) Blood 100,) Nat Struct Mol Bio W. (2013) Exp Biol rong, S.A. (2013) Bl | i U S A 98, 12902-7. / 20, 341-8. <i>ell Biochem</i> 95, 429-36. <i>Chem</i> 280, 41725-31. 3419-28. 587-97. 0) <i>Dev Biol</i> 339, 240-9. 1-72. 397-409. 3710-8. <i>ol</i> 13, 713-9. <i>Med (Maywood)</i> 238, 315 <i>ood</i> 121, 4847-53. | -23 <i>.</i> | |

| Species Reactivity | Species reactivity is determined by testing in at least one approved application (e.g., western blot). | | |
|------------------------|---|--|--|
| Western Blot Buffer | IMPORTANT: For western blots, incubate membrane with diluted primary antibody in 5% w/v BSA, 1X TBS, 0.1% Tween® 20 at 4°C with gentle shaking, overnight. | | |
| Applications Key | W: Western Blotting IP: Immunoprecipitation IF-IC: Immunofluorescence (Immunocytochemistry) | | |
| Cross-Reactivity Key | H: Human M: Mouse R: Rat Mk: Monkey | | |
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