## Di-Methyl-Histone H3 (Lys4) Antibody



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## For Research Use Only. Not for Use in Diagnostic Procedures.

<b>Applications:</b> W, IP, IHC-P, IF-IC, ChIP	Reactivity: H M R Mk	<b>Sensitivity:</b> Endogenous	<b>MW (kDa):</b> 17	<b>Source/Isotype:</b> Rabbit	<b>UniProt ID:</b> #P68431	Entrez-Gene Id: 8350
Product Usage Information		Application Western Blotting Immunoprecipitation Immunohistochemistry (Paraffin) Immunofluorescence (Immunocytochemistry) Chromatin IP				<b>Dilution</b> 1:1000 1:25 1:300 1:400 1:25
Storage		Supplied in 10 mM sodium HEPES (pH 7.5), 150 mM NaCl, 100 $\mu$ g/ml BSA and 50% glycerol. Store at – 20°C. Do not aliquot the antibody.				
Specificity/Sensitivity		Di-Methyl-Histone H3 (Lys4) Antibody detects endogenous levels of histone H3 when di-methylated on Lys4. The antibody shows very slight cross-reactivity with mono-methylated Lys4, but does not cross-react with non-methylated or tri-methylated Lys4. In addition, the antibody does not cross-react with mono-methylated, di-methylated or tri-methylated histone H3 at Lys9, Lys27, Lys36 or histone H4 at Lys20.				
Species predicted to react Xenopus, Zebrafish based on 100% sequence homology						
Source / Purific	cation		are produced by immunizing animals with a synthetic peptide corresponding to f histone H3 in which Lys4 is di-methylated. Antibodies are purified by peptide hy.			
Background		The nucleosome, made up of four core histone proteins (H2A, H2B, H3, and H4), is the primary building				

block of chromatin. Originally thought to function as a static scaffold for DNA packaging, histones have now been shown to be dynamic proteins, undergoing multiple types of post-translational modifications, including acetylation, phosphorylation, methylation, and ubiquitination (1). Histone methylation is a major determinant for the formation of active and inactive regions of the genome and is crucial for the proper programming of the genome during development (2,3). Arginine methylation of histones H3 (Arg2, 17, 26) and H4 (Arg3) promotes transcriptional activation and is mediated by a family of protein arginine methyltransferases (PRMTs), including the co-activators PRMT1 and CARM1 (PRMT4) (4). In contrast, a more diverse set of histone lysine methyltransferases has been identified, all but one of which contain a conserved catalytic SET domain originally identified in the Drosophila Su(var)3-9, Enhancer of zeste, and Trithorax proteins. Lysine methylation occurs primarily on histones H3 (Lys4, 9, 27, 36, 79) and H4 (Lys20) and has been implicated in both transcriptional activation and silencing (4). Methylation of these lysine residues coordinates the recruitment of chromatin modifying enzymes containing methyl-lysine binding modules such as chromodomains (HP1, PRC1), PHD fingers (BPTF, ING2), tudor domains (53BP1), and WD-40 domains (WDR5) (5-8). The discovery of histone demethylases, such as PADI4, LSD1, JMJD1, JMJD2, and JHDM1, has shown that methylation is a reversible epigenetic marker (9).

## **Background References**

- 1. Peterson, C.L. and Laniel, M.A. (2004) Curr Biol 14, R546-51.
- 2. Kubicek, S. et al. (2006) Ernst Schering Res Found Workshop, 1-27.
- 3. Lin, W. and Dent, S.Y. (2006) Curr Opin Genet Dev 16, 137-42.
- 4. Lee, D.Y. et al. (2005) Endocr Rev 26, 147-70.
- 5. Daniel, J.A. et al. (2005) Cell Cycle 4, 919-26.
- 6. Shi, X. et al. (2006) Nature 442, 96-9.
- 7. Wysocka, J. et al. (2006) *Nature* 442, 86-90.
- 8. Wysocka, J. et al. (2005) *Cell* 121, 859-72.
- 9. Trojer, P. and Reinberg, D. (2006) Cell 125, 213-7.

**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 IHC-P: Immunohistochemistry (Paraffin) IF-IC:

Immunofluorescence (Immunocytochemistry) ChIP: Chromatin IP

Cross-Reactivity Key H: Human M: Mouse R: Rat Mk: Monkey

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