760

Tri-Methyl-Histone H3 (Lys79) Antibody



Orders: 877-616-CELL (2355)

orders@cellsignal.com

Support: 877-678-TECH (8324)

Web: info@cellsignal.com

cellsignal.com

3 Trask Lane | Danvers | Massachusetts | 01923 | USA

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Applications:	Reactivity: H M R Mk	Sensitivity: Endogenous	MW (kDa): 17	Source/Isotype: Rabbit	UniProt ID: #P68431	Entrez-Gene Id: 8350
Product Usage Information		Application Western Blotting			Dilution 1:1000	
Storage		Supplied in 10 mM sodium HEPES (pH 7.5), 150 mM NaCl, 100 μ g/ml BSA and 50% glycerol. Store at – 20°C. Do not aliquot the antibody.				
Specificity/Sensitivity		Tri-Methyl-Histone H3 (Lys79) Antibody recognizes endogenous levels of histone H3 protein only when tri-methylated at Lys79. This antibody may also show slight cross-reactivity toward histone H3 when dimethylated at Lys79. This antibody does not cross-react with methylated histone H3 (Lys4, 9, 27, or 36).				
Species predicted to react based on 100% sequence homology		D. melanogaster, Xeno	opus, Zebrafish, Ho	rse		
Source / Purification		Polyclonal antibodies are produced by immunizing animals with a synthetic peptide corresponding to residues surrounding tri-methyl-Lys79 of human histone H3 protein. Antibodies are purified by protein A and peptide affinity chromatography.				
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 <i>Drosophila</i> 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		 Peterson, C.L. and Laniel, M.A. (2004) Curr Biol 14, R546-51. Kubicek, S. et al. (2006) Ernst Schering Res Found Workshop, 1-27. Lin, W. and Dent, S.Y. (2006) Curr Opin Genet Dev 16, 137-42. Lee, D.Y. et al. (2005) Endocr Rev 26, 147-70. Daniel, J.A. et al. (2005) Cell Cycle 4, 919-26. Shi, X. et al. (2006) Nature 442, 96-9. Wysocka, J. et al. (2006) Nature 442, 86-90. Wysocka, J. et al. (2005) Cell 121, 859-72. 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

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

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