HDAC2 (3F3) Mouse mAb



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Applications: WB, IP, IF-IC	Reactivity:	Sensitivity: Endogenous	MW (kDa): 60	Source/Isotype: Mouse IgG1	UniProt ID: #Q92769	Entrez-Gene Id	
		Endogenous		kappa	"Q32103		
Product Usage Information	Ар	plication				Dilution	
	We	stern Blotting				1:1000	
	Imr	Immunoprecipitation				1:50	
	Imr	Immunofluorescence (Immunocytochemistry)				1:400	
Storage	•	Supplied in 10 mM sodium HEPES (pH 7.5), 150 mM NaCl, 100 μ g/ml BSA, 50% glycerol and less than 0.02% sodium azide. Store at -20 °C. Do not aliquot the antibody.					
Specificity / Sensitiv	,	HDAC2 (3F3) Mouse mAb detects endogenous levels of HDAC2 protein. The antiboder react with other HDAC proteins.					
Source / Purification	-	oclonal antibody is oxy terminus of hur	responding to the				
Background		Acetylation of the histone tail causes chromatin to adopt an "open" conformation, allowing increased accessibility of transcription factors to DNA. The identification of histone acetyltransferases (HATs) and					

accessibility of transcription factors to DNA. The identification of histone acetyltransferases (HATs) and their large multiprotein complexes has yielded important insights into how these enzymes regulate transcription (1,2). HAT complexes interact with sequence-specific activator proteins to target specific genes. In addition to histones, HATs can acetylate nonhistone proteins, suggesting multiple roles for these enzymes (3). In contrast, histone deacetylation promotes a "closed" chromatin conformation and typically leads to repression of gene activity (4). Mammalian histone deacetylases can be divided into three classes on the basis of their similarity to various yeast deacetylases (5). Class I proteins (HDACs 1, 2, 3, and 8) are related to the yeast Rpd3-like proteins, those in class II (HDACs 4, 5, 6, 7, 9, and 10) are related to yeast Hda1-like proteins, and class III proteins are related to the yeast protein Sir2. Inhibitors of HDAC activity are now being explored as potential therapeutic cancer agents (6,7).

HDAC1 and HDAC2 are highly homologous and are involved in histone deacetylation, chromatin remodeling and transcriptional repression (8-10). Both proteins are found together in numerous complexes including the nucleosome remodeling and deacetylation complex (NuRD), MeCP1, and the mSin3A corepressor complex.

Background References

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- 2. Gregory, P.D. et al. (2001) Exp Cell Res 265, 195-202.
- 3. Liu, Y. et al. (2000) Mol Cell Biol 20, 5540-53.
- 4. Cress, W.D. and Seto, E. (2000) J Cell Physiol 184, 1-16.
- 5. Gray, S.G. and Ekström, T.J. (2001) *Exp Cell Res* 262, 75-83.
- 6. Thiagalingam, S. et al. (2003) Ann. N.Y. Acad. Sci. 983, 84-100.
- 7. Vigushin, D.M. and Coombes, R.C. (2004) Curr Cancer Drug Targets 4, 205-18.
- 8. Zhang, Y. et al. (1999) Genes Dev 13, 1924-35.
- 9. Ng, H.H. et al. (1999) Nat Genet 23, 58-61.
- 10. Zhang, Y. et al. (1997) Cell 89, 357-64.

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 nonfat dry milk, 1X TBS, 0.1% Tween® 20 at 4°C with gentle shaking, overnight.

Applications Key

Cross-Reactivity Key

WB: Western Blotting IP: Immunoprecipitation IF-IC: Immunofluorescence (Immunocytochemistry)

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H: human M: mouse R: rat Hm: hamster Mk: monkey Vir: virus Mi: mink C: chicken Dm: D. melanogaster X: Xenopus Z: zebrafish B: bovine Dq: dog Pq: pig Sc: S. cerevisiae Ce: C. elegans Hr: horse

GP: Guinea Pig **Rab:** rabbit **All:** all species expected

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