

## Acetyl-Histone H3-K4 Rabbit mAb

Catalog No.: RA8014

Basic Information	Background
<b>Observed MW</b> 17KDa	Histones are the main constituents of the protein part of chromosomes of eukaryotic cells. They are rich in the
Calculated MW 16kDa	amino acids arginine and lysine and have been greatly conserved during evolution. Histones pack the DNA into tight masses of chromatin. Two core histones of each
Category	class H2A, H2B, H3 and H4 assemble and are wrapped by
Primary antibody	146 base pairs of DNA to form one octameric nucleosome. Histone tails undergo numerous post-
Applications	translational modifications, which either directly or
IF/ICC,ChIP	indirectly alter chromatin structure to facilitate transcriptional activation or repression or other nuclear
Cross-Reactivity	processes. In addition to the genetic code, combinations
Human Yeast	of the different historie modifications reveal the so- called "histone code". Histone methylation and demethylation is dynamically regulated by respectively histone methyl transferases and histone demethylases.

## **Recommended Dilutions**

ELISA	1: 500	Source	Rabbit
WB	1: 500	Isotype	lgG
Dot Blot	1: 10000	Purification	Affinity purified.
IF/ICC	1: 300	Storage Conditions	Store at -20°C; for long storage, store at -80°C. Avoid multiple freeze-thaw cycles.
ChIP	0.5 µg	Storage buffer	In PBS with 0.05% sodium azide and 0.05% ProClin 300.

**Product Information** 



Note: For in vitro research use only, not for diagnostic or therapeutic use, This product is not a medical device. 注意:在体外研究使用,不用于诊断或治疗用途,本产品不是医疗装置!





Wild type and H3K4R mutant S. pombe cells were stained with both the antibody (in red) and by Hoechst staining (in blue, left), or with the antibody alone (right). The antibody was used at a dilution of 1:300.



HeLa cells extracts (15  $\mu\text{g})$  were analysed by WB blot using the antibody





ChIP assays were performed using WT and H3K4R mutant S. pombe and the antibody and optimized PCR primer sets for qPCR. The Fig shows the recovery, expressed as a % of input (the relative amount of IP DNA compared to input DNA after qPCR analysis).

To determine the titer, an ELISA was performed using a serial dilution of the antibody. The antigen used was a peptide containing the histone modification of interest. By plotting the absorbance against the antibody dilution the titer of the antibody was estimated to be 1:27800.

	H3K4un	H3K4me1	H3K4me2	H3K4me3	H3K4ac	H3K9ac	H3K14ac	H3K18ac	H3K23ac	H3K27ac	H3K36ac	H3K56ac
100 pmol												
50 pmol												
25 pmol												
5 pmol												
2 pmol					•							
0,2 pmol												
- ctri												

A Dot Blot analysis was performed to test the cross reactivity of the antibody with peptides containing other modifications of histone H3 and the unmodified H3K4 sequence. 100 to 0.2 pmol of the peptide containing the respective histone modification were spotted on a membrane. The Fig shows a high specificity of the antibody for the modification of interest.



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