

Histone H2B Recombinant Rabbit mAb

Catalog # AP98687

Product Information

Application	WB, IHC-P, IHC-F, IF, IP, CHIP, E
Primary Accession	P62807
Other Accession	P62807
Host	Rabbit
Clonality	Recombinant
Calculated MW	13906
Isotype	IgG
Purity	affinity purified by Protein A

Buffer 0.01M TBS (pH7.4) with 1% BSA, 0.02% Proclin300 and 50% Glycerol.

Background Descriptions Histones are basic nuclear proteins that are responsible for the nucleosome structure of the chromosomal fiber in eukaryotes. Nucleosomes consist of approximately 146 bp of DNA wrapped around a histone octamer composed of pairs of each of the four core histones (H2A, H2B, H3, and H4). The chromatin fiber is further compacted through the interaction of a linker histone, H1, with the DNA between the nucleosomes to form higher order chromatin structures. This gene is intronless and encodes a replication-dependent histone that is a member of the histone H2B family. Two transcripts that encode the same protein have been identified for this gene, which is found in the large histone gene cluster on chromosome 6p22-p21.3. [provided by RefSeq, Aug 2015],

Additional Information

Gene ID	3017;8339;8343;8344;8346;8347
Other Names	Histone H2B;Histone H2B type 1-C/E/F/G/I;Histone H2B.1 A;Histone H2B.a;H2B/a;Histone H2B.g;H2B/g;Histone H2B.h;H2B/h;Histone H2B.k;H2B/k;Histone H2B.l;H2B/l
Dilution	IHC-P=1:50-200,IHC-F=1:50-200,WB=1:500-2000,IF=1:50-200,ELISA=1:5000-10000,IP=1:50-200,ChIP=1:50-100
Storage	Store at -20 °C for one year. Avoid repeated freeze/thaw cycles. When reconstituted in sterile pH 7.4 0.01M PBS or diluent of antibody the antibody is stable for at least two weeks at 2-4 °C.

Protein Information

Name	H2BC4 (HGNC:4757)
Synonyms	H2BFL, HIST1H2BC

Function	Core component of nucleosome. Nucleosomes wrap and compact DNA into chromatin, limiting DNA accessibility to the cellular machineries which require DNA as a template. Histones thereby play a central role in transcription regulation, DNA repair, DNA replication and chromosomal stability. DNA accessibility is regulated via a complex set of post-translational modifications of histones, also called histone code, and nucleosome remodeling.
Cellular Location	Nucleus. Chromosome.

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