

Histone H2A (Acetyl Lys120) rabbit pAb

Catalog No: YK0134

Reactivity: Human; Mouse; Rat

Applications: WB;ELISA

Target: Histone H2A

Fields: >>Necroptosis;>>Neutrophil extracellular trap

formation;>>Alcoholism;>>Systemic lupus erythematosus

Gene Name: HIST1H2AG H2AFP; HIST1H2AI H2AFC; HIST1H2AK H2AFD; HIST1H2AL

H2AFI; HIST1H2AM H2AFN

Protein Name: Histone H2A (Acetyl Lys120)

Human Gene Id: 8329

Human Swiss Prot P0C0S8/P16104

No:

Mouse Gene ld: 319164

Mouse Swiss Prot

No:

Rat Swiss Prot No: P02262

Immunogen: Synthesized peptide derived from human Histone H2A (Acetyl Lys120)

Specificity: This antibody detects endogenous levels of Human, Mouse, Rat Histone H2A

(Acetyl Lys120)

P22752

Formulation: Liquid in PBS containing 50% glycerol, 0.5% BSA and 0.02% sodium azide.

Source: Polyclonal, Rabbit, lgG

Dilution: WB 1:1000-2000 ELISA 1:5000-20000

Purification: The antibody was affinity-purified from rabbit serum by affinity-chromatography



using specific immunogen.

Concentration: 1 mg/ml

Storage Stability: -15°C to -25°C/1 year(Do not lower than -25°C)

Observed Band: 15kD

Background: Histones are basic nuclear proteins that are responsible for the nucleosome

structure of the chromosomal fiber in eukaryotes. Two molecules of each of the four core histones (H2A, H2B, H3, and H4) form an octamer, around which approximately 146 bp of DNA is wrapped in repeating units, called nucleosomes. The linker histone, H1, interacts with linker DNA between nucleosomes and functions in the compaction of chromatin into higher order structures. This gene is intronless and encodes a replication-dependent histone that is a member of the histone H2A family. Transcripts from this gene lack polyA tails but instead contain a palindromic termination element. This gene is found in the small histone gene

cluster on chromosome 6p22-p21.3. [provided by RefSeq, Aug 2015],

Function: 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., mass

spectrometry: Monoisotopic with N-acetylserine

PubMed:16457589,PTM:Deiminated on Arg-4 in granulocytes upon calcium entry.,PTM:Monoubiquitination of Lys-120 by RING1 and RNF2/RING2 complex gives a specific tag for epigenetic transcriptional repression and participates in X chromosome inactivation of female mammals. It is involved in the initiation of both imprinted and random X inactivation. Ubiquitinated H2A is enriched in inactive X

chromosom

Subcellular Location:

Nucleus. Chromosome.

Products Images