

Histone H2A (Phospho Ser129) Polyclonal Antibody

Catalog No: YP1654

Reactivity: Human; Mouse; Rat

Applications: WB

Target: Histone H2A

Fields: >>Necroptosis;>>Neutrophil extracellular trap

formation;>>Alcoholism;>>Systemic lupus erythematosus

Gene Name: HIST1H2AG/HIST1H2AI/HIST1H2AK/HIST1H2AL/HIST1H2AM/HIST2H2AA3

/HIST2H2AA4/HIST3H2A

Protein Name: Histone H2A type 1/Histone H2A type 2/Histone H2A type 3

Human Gene Id: 8329/8330/8332/8336/8969/723790/8337/92815

Human Swiss Prot

No:

P0C0S8/Q6FI13/Q7L7L0

Mouse Gene ld: 319164/15267/319162

Rat Gene Id: 365877/64646

Rat Swiss Prot No: P02262/P0CC09/Q4FZT6

Immunogen: Synthetic Peptide of Histone H2A (Phospho Ser129)

Specificity: The antibody detects endogenous Histone H2A (Phospho Ser129) protein.

Formulation : PBS, pH 7.4, containing 0.5%BSA, 0.02% sodium azide as Preservative and

50% Glycerol.

Source : Polyclonal, Rabbit, IgG

Dilution: WB 1:1000-2000

Purification: The antibody was affinity-purified from rabbit antiserum by affinity-

1/3



chromatography using specific immunogen.

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

Observed Band: 14kD

Cell Pathway: Systemic lupus erythematosus;

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.

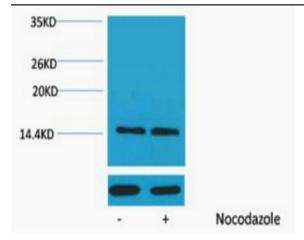
Expression : Bone, Brain, Colon, Eye, Lymph, PCR rescued clones, Placenta, Sple

Tag: orthogonal

Sort: 7426

No4:

Products Images



Western blot analysis of extracts from Hela cells, untreated (-) or treated, 1:5000. Secondary antibody(catalog#:RS0002) was diluted at 1:20000