

Histone H4 (Tri Methyl Lys59) Polyclonal Antibody

Catalog No: YH0023

Reactivity: Human; Mouse; Rat

Applications: WB

Target: Histone H4

Fields: >>Neutrophil extracellular trap formation;>>Alcoholism;>>Viral

carcinogenesis;>>Systemic lupus erythematosus

Gene Name: HIST1H4A H4/A H4FA; HIST1H4B H4/I H4FI; HIST1H4C H4/G H4FG;

HIST1H4D H4/B H4FB; HIST1H4E H4/J H4FJ; HIST1H4F H4/C H4FC; HIST1H4H H4/H H4FH; HIST1H4I H4/M H4FM; HIST1H4J H4/E H4FE; HIST1H4K H4/D H4FD; HIST1H4L H4/K H4FK; HIST2H4A H4/N H4F2 H4FN

HIST2H4; HIST2H4B H4/O H4FO; HIST4H4

Protein Name: Histone H4

Human Gene Id: 121504

Human Swiss Prot

No:

Mouse Gene ld: 100041230

Mouse Swiss Prot

No:

Rat Gene Id: 100360950

Rat Swiss Prot No: P62804

Immunogen: Synthetic Peptide of Histone H4 (Tri Methyl Lys59)

Specificity: The antibody detects endogenous Histone H4 (Tri Methyl Lys59) protein.

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

50% Glycerol.

P62805

P62806



Source: Polyclonal, Rabbit, IgG

Dilution: WB 1:500-1000

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

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 H4 family. Transcripts from this gene lack polyA tails but instead contain a palindromic termination element. This gene is found in the histone microcluster on

chromosome 6p21.33. [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.,PTM:Acetylation at Lys-6, Lys-9, Lys-13 and Lys-17 occurs in coding regions of the genome but

not in heterochromatin.,PTM:Citrullination at Arg-4 by PADI4 impairs

methylation.,PTM:Monomethylated, dimethylated or trimethylated at Lys-21. Monomethylation is performed by SET8. Trimethylation is performed by

SUV420H1 and SUV420H2 and induces gene silencing.,PTM:Monomethylation at Arg-4 by PRMT1 favors acetylation at Lys-9 and Lys-13. Demethylation is p

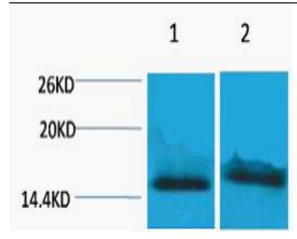
Subcellular Location:

Nucleus. Chromosome.

Expression:

B-cell lymphoma, Bone marrow, Brain, Clones donated by HIP, Corpus call

Products Images



Western blot analysis of 1) Hela, 2) 3T3, diluted at 1:2000. Secondary antibody(catalog#:RS0002) was diluted at 1:20000 cells nucleus extracted by Minute TM Cytoplasmic and Nuclear Fractionation kit (SC-003,Inventbiotech,MN,USA).