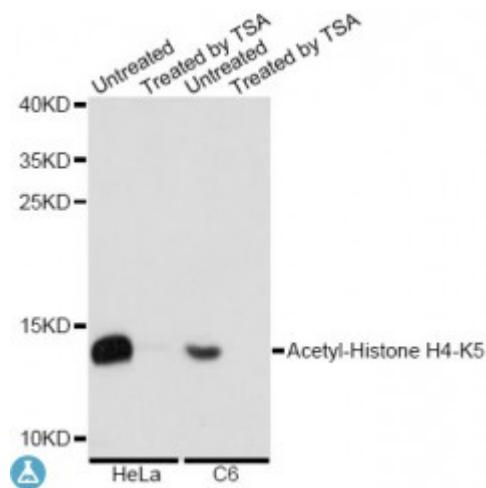


Anti-AcetylH4K5AC-(K5) Antibody



Description

Histones are basic nuclear proteins that are responsible for the nucleosome structure of the chromosomal fiber in eukaryotes. This structure consists of approximately 146 bp of DNA wrapped around a nucleosome, an 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 H4 family. Transcripts from this gene lack polyA tails; instead, they contain a palindromic termination element. This gene is found in a histone cluster on chromosome 1. This gene is one of four histone genes in the cluster that are duplicated; this record represents the centromeric copy.

| | |
|-----------------------|--|
| Model | STJ117427 |
| Host | Rabbit |
| Reactivity | Human, Rat |
| Applications | WB |
| Immunogen | A synthetic acetylated peptide around K5 of human Histone H4 (NP_001029249.1). |
| Gene ID | 121504 |
| Gene Symbol | HIST1H4A |
| Dilution range | WB 1:500 - 1:2000 |
| Purification | Affinity purification |
| Note | For Research Use Only (RUO). |

| | |
|---|---|
| Protein Name | Histone H4 |
| Molecular Weight | 11.367 kDa |
| Clonality | Monoclonal |
| Conjugation | Unconjugated |
| Isotype | IgG |
| Formulation | PBS with 0.02% sodium azide, 50% glycerol, pH7.3. |
| Storage Instruction | Store at -20C. Avoid freeze / thaw cycles. |
| Database Links | HGNC:4781 OMIM:142750 Reactome:R-HSA-1221632 |
| Alternative Names | Histone H4 |
| 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 Localization | Nucleus, Chromosome |
| Post-translational Modifications | Acetylation at Lys-6 (H4K5ac), Lys-9 (H4K8ac), Lys-13 (H4K12ac) and Lys-17 (H4K16ac) occurs in coding regions of the genome but not in heterochromatin, |

St John's Laboratory Ltd

F +44 (0)207 681 2580
T +44 (0)208 223 3081

W <http://www.stjohnslabs.com/>
E info@stjohnslabs.com