# abcam

# Product datasheet

# Anti-Histone H4 (mono methyl K31) antibody [EPR17913] ab177822

Recombinant

RabMAb

# 4 Images

#### Overview

Product name Anti-Histone H4 (mono methyl K31) antibody [EPR17913]

**Description** Rabbit monoclonal [EPR17913] to Histone H4 (mono methyl K31)

Host species Rabbit

Tested applications Suitable for: WB, ICC/IF, PepArr

Species reactivity Reacts with: Human

**Immunogen** Synthetic peptide. This information is proprietary to Abcam and/or its suppliers.

Positive control WB: HeLa whole cell lysate. ICC/IF: HeLa cells.

General notes This product is a recombinant monoclonal antibody, which offers several advantages including:

- High batch-to-batch consistency and reproducibility

- Improved sensitivity and specificity

- Long-term security of supply

- Animal-free production

For more information see here.

Our RabMAb<sup>®</sup> technology is a patented hybridoma-based technology for making rabbit monoclonal antibodies. For details on our patents, please refer to **RabMAb**<sup>®</sup> **patents**.

# **Properties**

Form Liquid

**Storage instructions** Shipped at 4°C. Store at +4°C short term (1-2 weeks). Upon delivery aliquot. Store at -20°C long

term. Avoid freeze / thaw cycle.

Storage buffer Preservative: 0.01% Sodium azide

Constituents: 59% PBS, 40% Glycerol (glycerin, glycerine), 0.05% BSA

Purity Protein A purified

ClonalityMonoclonalClone numberEPR17913

**Isotype** IgG

1

#### **Applications**

#### The Abpromise quarantee

Our **Abpromise guarantee** covers the use of ab177822 in the following tested applications.

The application notes include recommended starting dilutions; optimal dilutions/concentrations should be determined by the end user.

Application	Abreviews	Notes
WB		1/1000. Detects a band of approximately 11 kDa (predicted molecular weight: 11 kDa).
ICC/IF		1/2000.
PepArr		Use at an assay dependent concentration.

#### **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.

#### Sequence similarities

# Post-translational modifications

Belongs to the histone H4 family.

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.

Citrullination at Arg-4 (H4R3ci) by PADI4 impairs methylation.

Monomethylation and asymmetric dimethylation at Arg-4 (H4R3me1 and H4R3me2a, respectively) by PRMT1 favors acetylation at Lys-9 (H4K8ac) and Lys-13 (H4K12ac).

Demethylation is performed by JMJD6. Symmetric dimethylation on Arg-4 (H4R3me2s) by the

PRDM1/PRMT5 complex may play a crucial role in the germ-cell lineage.

Monomethylated, dimethylated or trimethylated at Lys-21 (H4K20me1, H4K20me2, H4K20me3).

Monomethylation is performed by SET8. Trimethylation is performed by SUV420H1 and

SUV420H2 and induces gene silencing.

Ubiquitinated by the CUL4-DDB-RBX1 complex in response to ultraviolet irradiation. This may weaken the interaction between histones and DNA and facilitate DNA accessibility to repair proteins. Monoubiquitinated at Lys-92 of histone H4 (H4K91ub1) in response to DNA damage. The exact role of H4K91ub1 in DNA damage response is still unclear but it may function as a licensing signal for additional histone H4 post-translational modifications such as H4 Lys-21

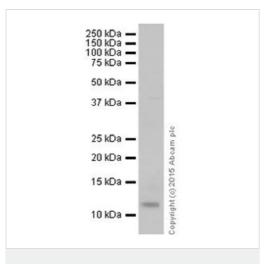
methylation (H4K20me).

Sumoylated, which is associated with transcriptional repression.

## **Cellular localization**

Nucleus. Chromosome.

### **Images**



Western blot - Anti-Histone H4 (mono methyl K31) antibody [EPR17913] (ab177822)

Anti-Histone H4 (mono methyl K31) antibody [EPR17913] (ab177822) at 1/1000 dilution + HeLa (Human epithelial cells from cervix adenocarcinoma) whole cell lysate at 10 μg

#### **Secondary**

Goat Anti-Rabbit IgG H&L (HRP) (ab97051) at 1/100000 dilution

**Predicted band size:** 11 kDa **Observed band size:** 11 kDa

Exposure time: 1 second

Blocking/Dilution buffer: 5% BSA/TBST.

ab177822 DAPI

-ve control 1

MERGED

-ve control 2

Immunocytochemistry/ Immunofluorescence - Anti-Histone H4 (mono methyl K31) antibody [EPR17913] (ab177822)

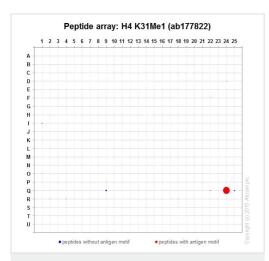
Immunofluorescent analysis of 4% paraformaldehyde-fixed, 0.1% Triton X-100 permeabilized HeLa (Human epithelial cells from cervix adenocarcinoma) cells labeling Histone H4 (mono methyl K31) with ab177822 at 1/2000 dilution, followed by Goat Anti-Rabbit IgG (Alexa Fluor<sup>®</sup> 488) (<u>ab150077</u>) secondary antibody at 1/1000 dilution (green). Confocal image showing nuclear staining on HeLa cell line. The nuclear counter stain is DAPI (blue).

Tubulin is detected with Anti-alpha Tubulin mouse MAb (ab7291) at 1/1000 dilution, followed by Goat Anti-Mouse IgG H&L (Alexa Fluor<sup>®</sup> 594) (ab150120) secondary antibody at 1/1000 dilution (red).

The negative controls are as follows:

-ve control 1: ab177822 at 1/2000 dilution, followed by Goat Anti-Mouse IgG H&L (Alexa Fluor<sup>®</sup> 594) (ab150120) secondary at 1/1000 dilution.

-ve control 2: Anti-alpha Tubulin mouse MAb ( $\underline{ab7291}$ ) at 1/1000 dilution, followed by Goat Anti-Rabbit lgG H&L (Alexa Fluor  $^{\mathbb{R}}$  488) ( $\underline{ab150077}$ ) secondary at 1/1000 dilution.

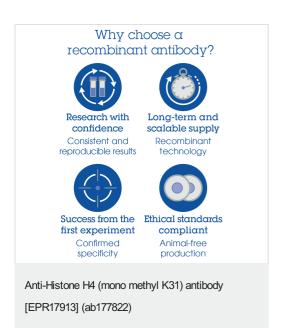


Peptide Array - Anti-Histone H4 (mono methyl K31) antibody [EPR17913] (ab177822)

ab177822 was tested in Peptide array against 501 different modified and unmodified histone peptides; each peptide is printed on the array at six concentrations (each in triplicate).

Circle area represents affinity between the antibody and a peptide: all antigen-containing peptides are displayed as red circles, all other peptides as blue circles. The affinity is calculated as area under curve when antibody binding values are plotted against the corresponding peptide concentration. Each circle area is normalized to the peptide with the strongest affinity.

The complete dataset, including full list of all peptides and information on the position of each peptide in the diagram, can be downloaded **here**.



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