abcam

Product datasheet

Anti-Histone H3 (acetyl K18) antibody - ChIP Grade ab1191

★★★★★ 8 Abreviews 179 References 画像数 6

製品の概要

製品名 Anti-Histone H3 (acetyl K18) antibody - ChIP Grade

製品の詳細 Rabbit polyclonal to Histone H3 (acetyl K18) - ChIP Grade

由来種 Rabbit

特異性 Specificity for acetyl K18 has been validated by blocking peptide/antibody incubation followed by

immunostaining western blots.

アプリケーション 適用あり: WB, PepArr, IHC-P, ICC/IF, ChIP

種交差性 交差種: Mouse, Human, Arabidopsis thaliana

交差が予測される動物種: Rat, Cow, Saccharomyces cerevisiae, Drosophila melanogaster,

Fungi 🕰

免疫原 Synthetic peptide. This information is proprietary to Abcam and/or its suppliers.

ポジティブ・コントロール ICC/IF: HeLa cells.

特記事項 The Life Science industry has been in the grips of a reproducibility crisis for a number of years.

Abcam is leading the way in addressing this with our range of recombinant monoclonal antibodies and knockout edited cell lines for gold-standard validation. Please check that this product meets

your needs before purchasing.

If you have any questions, special requirements or concerns, please send us an inquiry and/or contact our Support team ahead of purchase. Recommended alternatives for this product can be

found below, along with publications, customer reviews and Q&As

製品の特性

製品の状態 Liquid

保存方法 Shipped at 4°C. Store at +4°C short term (1-2 weeks). Upon delivery aliquot. Store at -20°C or -

80°C. Avoid freeze / thaw cycle.

バッファー pH: 7.40

Preservative: 0.02% Sodium azide

Constituent: PBS

Batches of this product that have a concentration < 1mg/ml may have BSA added as a stabilising

agent. If you would like information about the formulation of a specific lot, please contact our

scientific support team who will be happy to help.

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精製度 Immunogen affinity purified

ポリ/モノ ポリクローナル

アイソタイプ IgG

アプリケーション

The Abpromise guarantee Abpromise保証は、次のテスト済みアプリケーションにおけるab1191の使用に適用されますアプリケーションノートには、推奨の開始希釈率がありますが、適切な希釈率につきましてはご検討ください。

アプリケーション	Abreviews	特記事項
WB	★★★★★ (3)	1/1000. Detects a band of approximately 17 kDa.
PepArr		Use a concentration of 0.2 - 0.02 μg/ml.
IHC-P		1/200.
ICC/IF	**** (1)	Use a concentration of 1 µg/ml.
ChIP	★★★★ (4)	Use 2µg for 10 ⁶ cells.

ターゲット情報

機能

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.

配列類似性

発生段階

翻訳後修飾

Belongs to the histone H3 family.

Expressed during S phase, then expression strongly decreases as cell division slows down

during the process of differentiation.

Acetylation is generally linked to gene activation. Acetylation on Lys-10 (H3K9ac) impairs methylation at Arg-9 (H3R8me2s). Acetylation on Lys-19 (H3K18ac) and Lys-24 (H3K24ac) favors methylation at Arg-18 (H3R17me).

Citrullination at Arg-9 (H3R8ci) and/or Arg-18 (H3R17ci) by PAD4 impairs methylation and represses transcription.

Asymmetric dimethylation at Arg-18 (H3R17me2a) by CARM1 is linked to gene activation. Symmetric dimethylation at Arg-9 (H3R8me2s) by PRMT5 is linked to gene repression.

Asymmetric dimethylation at Arg-3 (H3R2me2a) by PRMT6 is linked to gene repression and is mutually exclusive with H3 Lys-5 methylation (H3K4me2 and H3K4me3). H3R2me2a is present at the 3' of genes regardless of their transcription state and is enriched on inactive promoters, while it is absent on active promoters.

Methylation at Lys-5 (H3K4me), Lys-37 (H3K36me) and Lys-80 (H3K79me) are linked to gene activation. Methylation at Lys-5 (H3K4me) facilitates subsequent acetylation of H3 and H4. Methylation at Lys-80 (H3K79me) is associated with DNA double-strand break (DSB) responses and is a specific target for TP53BP1. Methylation at Lys-10 (H3K9me) and Lys-28 (H3K27me) are linked to gene repression. Methylation at Lys-10 (H3K9me) is a specific target for HP1 proteins (CBX1, CBX3 and CBX5) and prevents subsequent phosphorylation at Ser-11

(H3S10ph) and acetylation of H3 and H4. Methylation at Lys-5 (H3K4me) and Lys-80 (H3K79me) require preliminary monoubiquitination of H2B at 'Lys-120'. Methylation at Lys-10 (H3K9me) and Lys-28 (H3K27me) are enriched in inactive X chromosome chromatin.

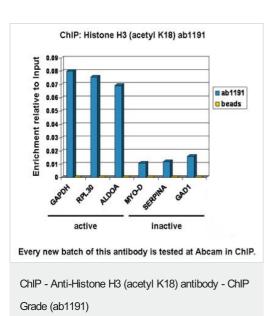
Phosphorylated at Thr-4 (H3T3ph) by GSG2/haspin during prophase and dephosphorylated during anaphase. Phosphorylation at Ser-11 (H3S10ph) by AURKB is crucial for chromosome condensation and cell-cycle progression during mitosis and meiosis. In addition phosphorylation at Ser-11 (H3S10ph) by RPS6KA4 and RPS6KA5 is important during interphase because it enables the transcription of genes following external stimulation, like mitogens, stress, growth factors or UV irradiation and result in the activation of genes, such as c-fos and c-jun. Phosphorylation at Ser-11 (H3S10ph), which is linked to gene activation, prevents methylation at Lys-10 (H3K9me) but facilitates acetylation of H3 and H4. Phosphorylation at Ser-11 (H3S10ph) by AURKB mediates the dissociation of HP1 proteins (CBX1, CBX3 and CBX5) from heterochromatin. Phosphorylation at Ser-11 (H3S10ph) is also an essential regulatory mechanism for neoplastic cell transformation. Phosphorylated at Ser-29 (H3S28ph) by MLTK isoform 1, RPS6KA5 or AURKB during mitosis or upon ultraviolet B irradiation. Phosphorylation at Thr-7 (H3T6ph) by PRKCBB is a specific tag for epigenetic transcriptional activation that prevents demethylation of Lys-5 (H3K4me) by LSD1/KDM1A. At centromeres, specifically phosphorylated at Thr-12 (H3T11ph) from prophase to early anaphase, by DAPK3 and PKN1. Phosphorylation at Thr-12 (H3T11ph) by PKN1 is a specific tag for epigenetic transcriptional activation that promotes demethylation of Lys-10 (H3K9me) by KDM4C/JMJD2C. Phosphorylation at Tyr-42 (H3Y41ph) by JAK2 promotes exclusion of CBX5 (HP1 alpha) from chromatin.

Monoubiquitinated by RAG1 in lymphoid cells, monoubiquitination is required for V(D)J recombination (By similarity). 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.

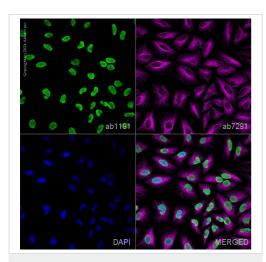
Nucleus. Chromosome.

細胞内局在

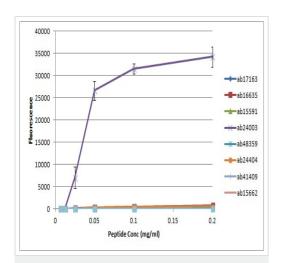
画像



Chromatin was prepared from U2OS cells according to the Abcam X-ChIP protocol. Cells were fixed with formaldehyde for 10 min. The ChIP was performed with 25 μ g of chromatin, 2 μ g of ab1191 (blue), and 20 μ l of protein A/G sepharose beads. No antibody was added to the beads control (yellow). The immunoprecipitated DNA was quantified by real time PCR (Taqman approach). Primers and probes are located in the first kb of the transcribed region.



Immunocytochemistry/ Immunofluorescence - Anti-Histone H3 (acetyl K18) antibody - ChIP Grade (ab1191)



Peptide Array - Anti-Histone H3 (acetyl K18) antibody - ChIP Grade (ab1191)

ab1191 staining Histone H3 (acetyl K18) in HeLa cells. The cells were fixed with 100% methanol (5 min), permeabilized with 0.1% PBS-Triton X-100 for 5 minutes and then blocked with 1% BSA/10% normal goat serum/0.3M glycine in 0.1%PBS-Tween for 1h. The cells were then incubated overnight at 4°C with ab1191 at 0.1µg/ml and ab7291, Mouse monoclonal [DM1A] to alpha Tubulin -Loading Control. Cells were then incubated with ab150081, Goat polyclonal Secondary Antibody to Rabbit IgG - H&L (Alexa Fluor® 488), pre-adsorbed at 1/1000 dilution (shown in green) and ab150120, Goat polyclonal Secondary Antibody to Mouse IgG -H&L (Alexa Fluor[®] 594), pre-adsorbed at 1/1000 dilution (shown in pseudocolour magenta). Nuclear DNA was labelled with DAPI (shown in blue). Also suitable in cells fixed with 4% paraformaldehyde (10 min).lmage was acquired with a high-content analyser (Operetta CLS, Perkin Elmer) and a maximum intensity projection of confocal sections is shown.

All batches of ab1191 are tested in Peptide Array against peptides to different Histone H3 and H4 modifications. Six dilutions of each peptide are printed on to the Peptide Array in triplicate and results are averaged before being plotted on to a graph. Results show strong binding to Histone H3 - acetyl K18 peptide (ab24003), indicating that this antibody specifically recognises the Histone H3 - acetyl K18 modification.

ab17163 - Histone H3 - unmodified

ab16635 - Histone H3 - acetyl K9

ab15591 - Histone H3 - acetyl K14

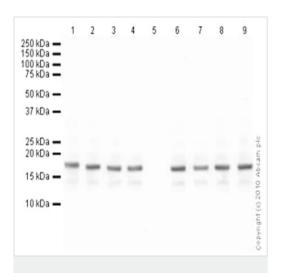
ab24003 - Histone H3 - acetyl K18

ab48359 - Histone H3 - acetyl K23

ab24404 - Histone H3 - acetyl K27

ab41409 - Histone H3 - acetyl K36

ab15662 - Histone H4 - acetyl K12



Western blot - Anti-Histone H3 (acetyl K18) antibody - ChIP Grade (ab1191)

All lanes : Anti-Histone H3 (acetyl K18) antibody - ChIP Grade (ab1191) at 1 µg/ml

Lane 1: HeLa Histone Preparation Nuclear Lysate

Lane 2: HeLa Histone Preparation Nuclear Lysate with Histone H3

peptide - unmodified R17 at 0.5 µg/ml

Lane 3: HeLa Histone Preparation Nuclear Lysate with Human

Histone H3 (acetyl K9) peptide (ab16635) at 0.5 µg/ml

Lane 4: HeLa Histone Preparation Nuclear Lysate with Histone H3

peptide - acetyl K14 at 0.5 µg/ml

Lane 5: HeLa Histone Preparation Nuclear Lysate with Human

Histone H3 (acetyl K18) peptide ($\underline{ab24003}$) at 0.5 $\mu g/ml$

Lane 6 : HeLa Histone Preparation Nuclear Lysate with Human

Histone H3 (acetyl K23) peptide (ab48359) at 0.5 µg/ml

Lane 7: HeLa Histone Preparation Nuclear Lysate with Human

Histone H3 (acetyl K27) peptide (ab24404) at 0.5 µg/ml

Lane 8: HeLa Histone Preparation Nuclear Lysate with Histone H3

peptide - acetyl K36 at 0.5 µg/ml

Lane 9: HeLa Histone Preparation Nuclear Lysate with Human

Histone H4 (acetyl K12) peptide (ab15662) at 0.5 µg/ml

Lysates/proteins at 2.5 µg per lane.

Secondary

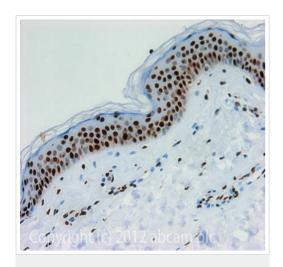
All lanes : Goat polyclonal to Rabbit lgG - H&L - Pre-Adsorbed (HRP) at 1/3000 dilution

Developed using the ECL technique.

Performed under reducing conditions.

Predicted band size: 15.4 kDa **Observed band size:** 17 kDa

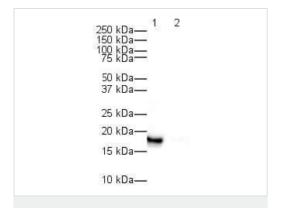
Exposure time: 1 minute



Immunohistochemistry (Formalin/PFA-fixed paraffinembedded sections) - Anti-Histone H3 (acetyl K18) antibody - ChIP Grade (ab1191)

IHC image of Histone H3 (acetyl K18) staining in Human normal skin formalin fixed paraffin embedded tissue section, performed on a Leica BondTM system using the standard protocol F. The section was pre-treated using heat mediated antigen retrieval with sodium citrate buffer (pH6, epitope retrieval solution 1) for 20 mins. The section was then incubated with ab1191, 1µg/ml, for 15 mins at room temperature and detected using an HRP conjugated compact polymer system. DAB was used as the chromogen. The section was then counterstained with haematoxylin and mounted with DPX.

For other IHC staining systems (automated and non-automated) customers should optimize variable parameters such as antigen retrieval conditions, primary antibody concentration and antibody incubation times.



Western blot - Anti-Histone H3 (acetyl K18) antibody - ChIP Grade (ab1191)

All lanes : Anti-Histone H3 (acetyl K18) antibody - ChIP Grade (ab1191) at 1 μg/ml

Lane 1: Histone prep

Lane 2 : Histone prep with Human Histone H3 (acetyl K18) peptide (ab24003) at 1 μ g/ml

Lysates/proteins at 0.5 µg/ml per lane.

Secondary

All lanes : Goat Anti-Rabbit lgG H&L (HRP) (ab6721) at 1/5000 dilution

Predicted band size: 15.4 kDa

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