abcam

Product datasheet

Anti-CTCF antibody [EPR7313] ab128909



RabMAb

3 Images

Overview

Product name Anti-CTCF antibody [EPR7313]

Description Rabbit monoclonal [EPR7313] to CTCF

Host species Rabbit

Tested applications Suitable for: WB

Unsuitable for: Flow Cyt,ICC/IF,IHC-P or IP

Species reactivity Reacts with: Mouse, Human

Predicted to work with: Rat ...

Immunogen Synthetic peptide within Human CTCF. The exact sequence is proprietary.

Positive control U937, HeLa, MDA-MB-435, 293T, Caco-2 and RAW 264.7 cell lysates.

General notesThis 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.

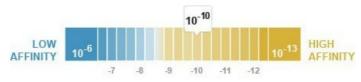
Our RabMAb[®] technology is a patented hybridoma-based technology for making rabbit monoclonal antibodies. For details on our patents, please refer to **RabMAb**[®] **patents**.

Properties

Form Liquid

Storage instructions Shipped at 4°C. Store at -20°C. Stable for 12 months at -20°C.

Dissociation constant (K_D) $K_D = 1.58 \times 10^{-10} M$



Learn more about K_D

Storage buffer pH: 7.20

Preservative: 0.01% Sodium azide

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Constituents: 9% PBS, 40% Glycerol (glycerin, glycerine), 0.05% BSA, 50% Tissue culture

supernatant

Purity Protein A purified

Clone number Monoclonal EPR7313

Isotype IgG

Applications

The Abpromise guarantee

Our **Abpromise guarantee** covers the use of ab128909 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. Predicted molecular weight: 83 kDa.

Application notes

Is unsuitable for Flow Cyt,ICC/IF,IHC-P or IP.

Target

Function

Chromatin binding factor that binds to DNA sequence specific sites. Involved in transcriptional regulation by binding to chromatin insulators and preventing interaction between promoter and nearby enhancers and silencers. Acts as transcriptional repressor binding to promoters of vertebrate MYC gene and BAG1 gene. Also binds to the PLK and PIM1 promoters. Acts as a transcriptional activator of APP. Regulates APOA1/C3/A4/A5 gene cluster and controls MHC class II gene expression. Plays an essential role in oocyte and preimplantation embryo development by activating or repressing transcription. Seems to act as tumor suppressor. Plays a critical role in the epigenetic regulation. Participates to the allele-specific gene expression at the imprinted IGF2/H19 gene locus. On the maternal allele, binding within the H19 imprinting control region (ICR) mediates maternally inherited higher-order chromatin conformation to restrict enhancer access to IGF2. Plays a critical role in gene silencing over considerable distances in the genome. Preferentially interacts with unmethylated DNA, preventing spreading of CpG methylation and maintaining methylation-free zones. Inversely, binding to target sites is prevented by CpG methylation. Plays a important role in chromatin remodeling. Can dimerize when it is bound to different DNA sequences, mediating long-range chromatin looping. Mediates interchromosomal association between IGF2/H19 and WSB1/NF1 and may direct distant DNA segments to a common transcription factory. Causes local loss of histone acetylation and gain of histone methylation in the beta-globin locus, without affecting transcription. When bound to chromatin, it provides an anchor point for nucleosomes positioning. Seems to be essential for homologous X-chromosome pairing. May participate with Tsix in establishing a regulatable epigenetic switch for X chromosome inactivation. May play a role in preventing the propagation of stable methylation at the escape genes from X- inactivation. Involved in sister chromatid cohesion. Associates with both centromeres and chromosomal arms during metaphase and required for cohesin localization to CTCF sites. Regulates asynchronous replication of IGF2/H19.

Tissue specificity

Ubiquitous. Absent in primary spermatocytes.

Sequence similarities

Belongs to the CTCF zinc-finger protein family.

Contains 11 C2H2-type zinc fingers.

Domain

The 11 zinc fingers are highly conserved among vertebrates, exhibiting almost identical amino acid sequences. Different subsets or combination of individual zinc fingers gives the ability to CTCF to recognize multiple DNA target sites.

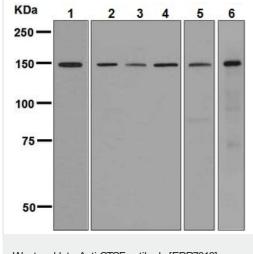
Post-translational modifications

Sumoylated on Lys-74 and Lys-689; sumoylation of CTCF contributes to the repressive function of CTCF on the MYC P2 promoter.

Cellular localization

Nucleus > nucleoplasm. Chromosome. Chromosome > centromere. May translocate to the nucleolus upon cell differentiation. Associates with both centromeres and chromosomal arms during metaphase. Associates with the H19 ICR in mitotic chromosomes. May be preferentially excluded from heterochromatin during interphase.

Images



All lanes : Anti-CTCF antibody [EPR7313] (ab128909) at 1/1000 dilution

Lane 1 : U937 cell lysate

Lane 2 : HeLa cell lysate

Lane 3: MDA-MB-435 cell lysate

Lane 4 : 293T cell lysate

Lane 5 : Caco-2 cell lysate

Lane 6 : RAW 264.7 cell lysate

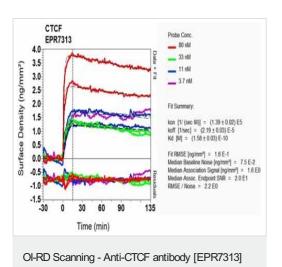
Lysates/proteins at 10 µg per lane.

Western blot - Anti-CTCF antibody [EPR7313] (ab128909)

Secondary

All lanes: HRP labelled goat anti-rabbit at 1/2000 dilution

Predicted band size: 83 kDa

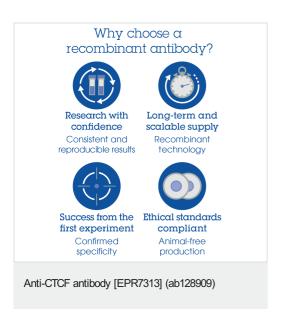


(ab128909)

Equilibrium disassociation constant (K_D)

Learn more about K_D

Click here to learn more about KD



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