

## Anti-CTCF Rabbit Monoclonal Antibody Catalog # ABO13523

### Specification

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#### Anti-CTCF Rabbit Monoclonal Antibody - Product Information

Application	WB, IHC, IF, ICC, FC
Primary Accession	<a href="#">P49711</a>
Host	Rabbit
Isotype	Rabbit IgG
Reactivity	Rat, Human, Mouse
Clonality	Monoclonal
Format	Liquid

#### Description

Anti-CTCF Rabbit Monoclonal Antibody . Tested in WB, IHC, ICC/IF, Flow Cytometry applications. This antibody reacts with Human, Mouse, Rat.

#### Anti-CTCF Rabbit Monoclonal Antibody - Additional Information

**Gene ID** 10664

#### Other Names

Transcriptional repressor CTCF, 11-zinc finger protein, CCCTC-binding factor, CTCFL paralog, CTCF

#### Calculated MW

82785 MW KDa

#### Application Details

WB 1:500-1:2000<br>IHC 1:100-1:500<br>ICC/IF 1:50-1:200<br>FC 1:50

#### Subcellular 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.

#### Tissue Specificity

Ubiquitous. Absent in primary spermatocytes..

#### Contents

Rabbit IgG in phosphate buffered saline, pH 7.4, 150mM NaCl, 0.02% sodium azide and 50% glycerol, 0.4-0.5mg/ml BSA.

#### Immunogen

A synthesized peptide derived from human CTCF

#### Purification

Affinity-chromatography

Storage

Store at -20°C for one year. For short term

storage and frequent use, store at 4°C for up to one month. Avoid repeated freeze-thaw cycles.

## Anti-CTCF Rabbit Monoclonal Antibody - Protein Information

### Name CTCF

### Function

Chromatin binding factor that binds to DNA sequence specific sites and regulates the 3D structure of chromatin (PubMed: [18347100](http://www.uniprot.org/citations/18347100)), PubMed: [18654629](http://www.uniprot.org/citations/18654629), PubMed: [19322193](http://www.uniprot.org/citations/19322193)). Binds together strands of DNA, thus forming chromatin loops, and anchors DNA to cellular structures, such as the nuclear lamina (PubMed: [18347100](http://www.uniprot.org/citations/18347100), PubMed: [18654629](http://www.uniprot.org/citations/18654629), PubMed: [19322193](http://www.uniprot.org/citations/19322193)). Defines the boundaries between active and heterochromatic DNA via binding to chromatin insulators, thereby preventing interaction between promoter and nearby enhancers and silencers (PubMed: [18347100](http://www.uniprot.org/citations/18347100), PubMed: [18654629](http://www.uniprot.org/citations/18654629), PubMed: [19322193](http://www.uniprot.org/citations/19322193)). Plays a critical role in the epigenetic regulation (PubMed: [16949368](http://www.uniprot.org/citations/16949368)). Participates in the allele-specific gene expression at the imprinted IGF2/H19 gene locus (PubMed: [16107875](http://www.uniprot.org/citations/16107875), PubMed: [16815976](http://www.uniprot.org/citations/16815976), PubMed: [17827499](http://www.uniprot.org/citations/17827499)). 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 (By similarity). Mediates interchromosomal association between IGF2/H19 and WSB1/NF1 and may direct distant DNA segments to a common transcription factory (By similarity). Regulates asynchronous replication of IGF2/H19 (By similarity). Plays a critical role in gene silencing over considerable distances in the genome (By similarity). Preferentially interacts with unmethylated DNA, preventing spreading of CpG methylation and maintaining methylation-free zones (PubMed: [18413740](http://www.uniprot.org/citations/18413740)). Inversely, binding to target sites is prevented by CpG methylation (PubMed: [18413740](http://www.uniprot.org/citations/18413740)). Plays an important role in chromatin remodeling (PubMed: [18413740](http://www.uniprot.org/citations/18413740)). Can dimerize when it is bound to different DNA sequences, mediating long-range chromatin looping (PubMed: [12191639](http://www.uniprot.org/citations/12191639)). Causes local loss of histone acetylation and gain of histone methylation in the beta-globin locus, without affecting transcription (PubMed: [12191639](http://www.uniprot.org/citations/12191639)). When bound to chromatin, it provides an anchor point for nucleosomes positioning (PubMed: [12191639](http://www.uniprot.org/citations/12191639)). Seems to be essential for homologous X-chromosome pairing (By similarity). May participate with Tsix in establishing a regulatable epigenetic switch for X chromosome inactivation (PubMed: [11743158](http://www.uniprot.org/citations/11743158)). May play a role in preventing the propagation of stable methylation at the escape genes from X-inactivation (PubMed: [11743158](http://www.uniprot.org/citations/11743158)). Involved in sister chromatid cohesion (PubMed: [12191639](http://www.uniprot.org/citations/12191639)). Associates with both centromeres and chromosomal arms during metaphase and required for cohesin localization to CTCF sites (PubMed: [12191639](http://www.uniprot.org/citations/12191639)).

<http://www.uniprot.org/citations/18550811> target="\_blank">18550811</a>). Plays a role in the recruitment of CENPE to the pericentromeric/centromeric regions of the chromosome during mitosis (PubMed:<a href="http://www.uniprot.org/citations/26321640" target="\_blank">26321640</a>). Acts as a transcriptional repressor binding to promoters of vertebrate MYC gene and BAG1 gene (PubMed:<a href="http://www.uniprot.org/citations/18413740" target="\_blank">18413740</a>, PubMed:<a href="http://www.uniprot.org/citations/8649389" target="\_blank">8649389</a>, PubMed:<a href="http://www.uniprot.org/citations/9591631" target="\_blank">9591631</a>). Also binds to the PLK and PIM1 promoters (PubMed:<a href="http://www.uniprot.org/citations/12191639" target="\_blank">12191639</a>). Acts as a transcriptional activator of APP (PubMed:<a href="http://www.uniprot.org/citations/9407128" target="\_blank">9407128</a>). Regulates APOA1/C3/A4/A5 gene cluster and controls MHC class II gene expression (PubMed:<a href="http://www.uniprot.org/citations/18347100" target="\_blank">18347100</a>, PubMed:<a href="http://www.uniprot.org/citations/19322193" target="\_blank">19322193</a>). Plays an essential role in oocyte and preimplantation embryo development by activating or repressing transcription (By similarity). Seems to act as tumor suppressor (PubMed:<a href="http://www.uniprot.org/citations/12191639" target="\_blank">12191639</a>).

### Cellular Location

Nucleus, nucleoplasm. Chromosome. Chromosome, centromere. Note=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

### Tissue Location

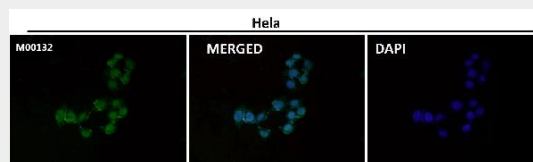
Ubiquitous. Absent in primary spermatocytes.

## Anti-CTCF Rabbit Monoclonal Antibody - Protocols

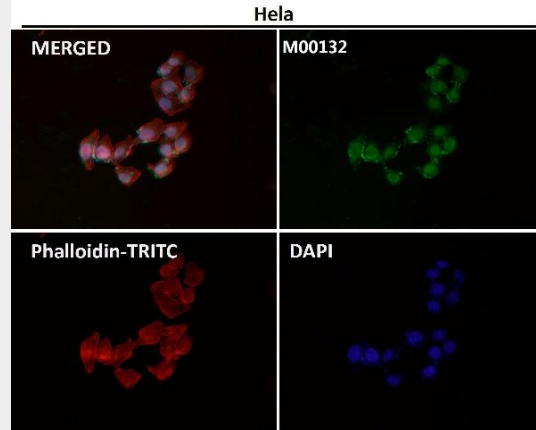
Provided below are standard protocols that you may find useful for product applications.

- [Western Blot](#)
- [Blocking Peptides](#)
- [Dot Blot](#)
- [Immunohistochemistry](#)
- [Immunofluorescence](#)
- [Immunoprecipitation](#)
- [Flow Cytometry](#)
- [Cell Culture](#)

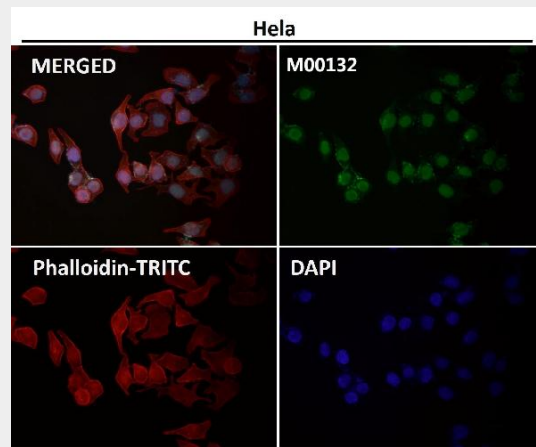
## Anti-CTCF Rabbit Monoclonal Antibody - Images



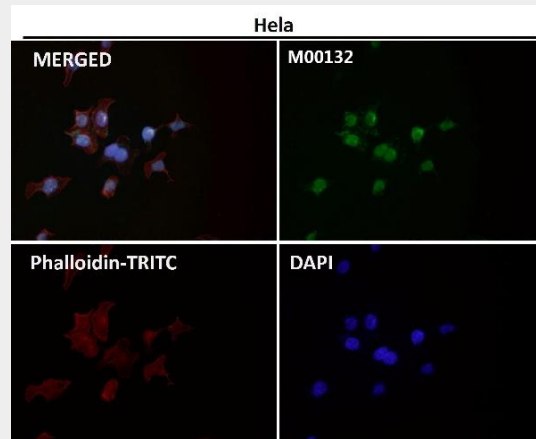
Immunofluorescent analysis using the Antibody at 1:50 dilution.



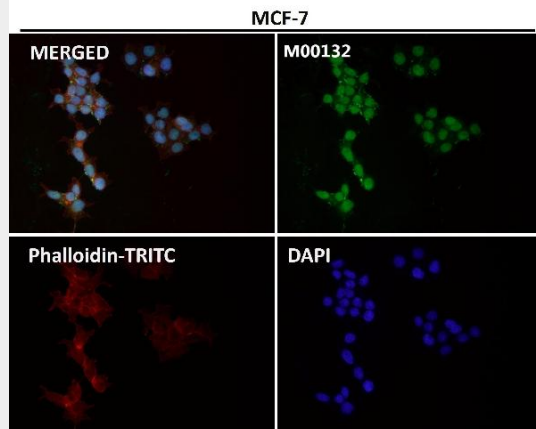
Immunofluorescent analysis using the Antibody at 1:50 dilution.



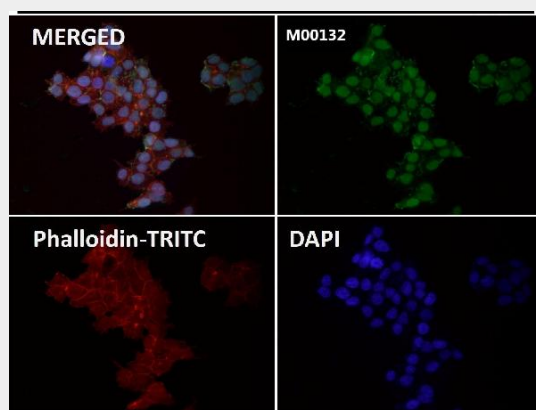
Immunofluorescent analysis using the Antibody at 1:150 dilution.



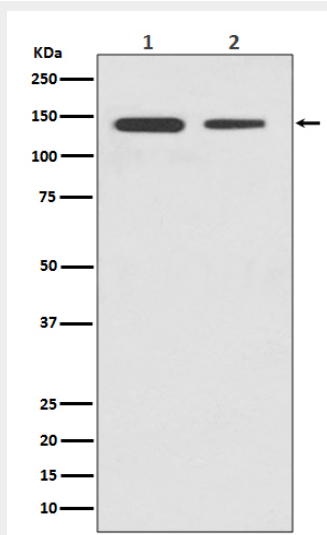
Immunofluorescent analysis using the Antibody at 1:500 dilution.



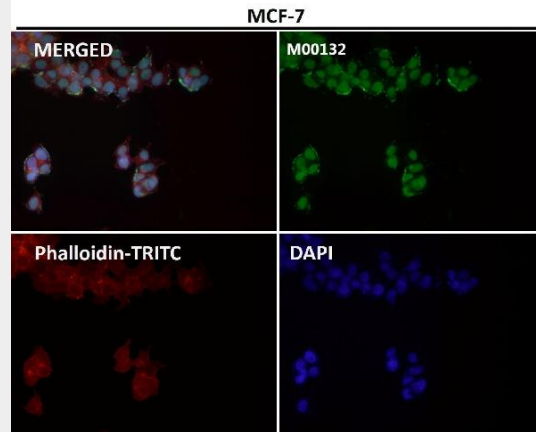
Immunofluorescent analysis using the Antibody at 1:50 dilution.



Immunofluorescent analysis using the Antibody at 1:150 dilution.



Western blot analysis of CTCF expression in (1) HeLa cell lysate; (2) Mouse brain lysate.



Immunofluorescent analysis using the Antibody at 1:500 dilution.