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CTR9 antibody - C-terminal region
Rabbit Polyclonal Antibody
Catalog # AI15141

Specification

CTR9 antibody - C-terminal region - Product Information

Application	**WB**
Primary Accession	Q6PD62
Other Accession	NM_014633, NP_055448
Reactivity	**Human, Mouse, Rat, Pig, Horse, Bovine, Guinea Pig, Dog**
Predicted	**Human, Mouse, Rat, Pig, Horse, Bovine, Guinea Pig, Dog**
Host	**Rabbit**
Clonality	**Polyclonal**
Calculated MW	**133kDa KDa**

CTR9 antibody - C-terminal region - Additional Information

Gene ID 9646

Alias Symbol **KIAA0155, SH2BP1, TSBP, p150, p150TSP**

Other Names
RNA polymerase-associated protein CTR9 homolog, SH2 domain-binding protein 1, CTR9, KIAA0155, SH2BP1

Format
Liquid. Purified antibody supplied in 1x PBS buffer with 0.09% (w/v) sodium azide and 2% sucrose.

Reconstitution & Storage
Add 50 ul of distilled water. Final anti-CTR9 antibody concentration is 1 mg/ml in PBS buffer with 2% sucrose. For longer periods of storage, store at 20°C. Avoid repeat freeze-thaw cycles.

Precautions
CTR9 antibody - C-terminal region is for research use only and not for use in diagnostic or therapeutic procedures.

CTR9 antibody - C-terminal region - Protein Information

Name CTR9

Synonyms KIAA0155, SH2BP1

Function
Component of the PAF1 complex (PAF1C) which has multiple functions during transcription by RNA polymerase II and is implicated in regulation of development and maintenance of embryonic stem cell pluripotency. PAF1C associates with RNA polymerase II through interaction with POLR2A CTD non-phosphorylated and 'Ser-2'- and 'Ser- 5'-phosphorylated forms and is involved in

transcriptional elongation, acting both independently and synergistically with TCEA1 and in cooperation with the DSIF complex and HTATSF1. PAF1C is required for transcription of Hox and Wnt target genes. PAF1C is involved in hematopoiesis and stimulates transcriptional activity of KMT2A/MLL1; it promotes leukemogenesis through association with KMT2A/MLL1-rearranged oncoproteins, such as KMT2A/MLL1-MLLT3/AF9 and KMT2A/MLL1-MLLT1/ENL. PAF1C is involved in histone modifications such as ubiquitination of histone H2B and methylation on histone H3 'Lys-4' (H3K4me3). PAF1C recruits the RNF20/40 E3 ubiquitin-protein ligase complex and the E2 enzyme UBE2A or UBE2B to chromatin which mediate monoubiquitination of 'Lys-120' of histone H2B (H2BK120ub1); UB2A/B-mediated H2B ubiquitination is proposed to be coupled to transcription. PAF1C is involved in mRNA 3' end formation probably through association with cleavage and poly(A) factors. In case of infection by influenza A strain H3N2, PAF1C associates with viral NS1 protein, thereby regulating gene transcription. Required for mono- and trimethylation on histone H3 'Lys-4' (H3K4me3) and dimethylation on histone H3 'Lys-79' (H3K4me3). Required for Hox gene transcription. Required for the trimethylation of histone H3 'Lys-4' (H3K4me3) on genes involved in stem cell pluripotency; this function is synergistic with CXXC1 indicative for an involvement of the SET1 complex. Involved in transcriptional regulation of IL6-responsive genes and in JAK-STAT pathway; may regulate DNA-association of STAT3 (By similarity).

Cellular Location
Nucleus speckle {ECO:0000250|UniProtKB:Q62018}.

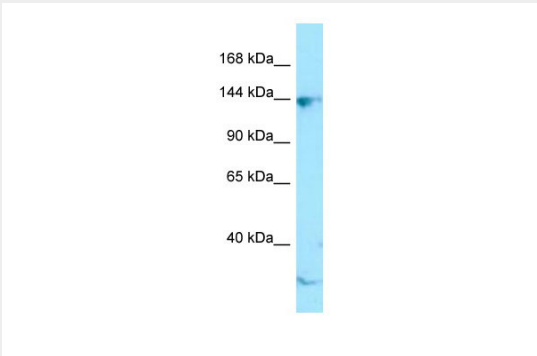
Tissue Location
Widely expressed..

CTR9 antibody - C-terminal region - Protocols

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

- Western Blot
- Blocking Peptides
- Dot Blot
- Immunohistochemistry
- Immunofluorescence
- Immunoprecipitation
- Flow Cytomety
- Cell Culture

CTR9 antibody - C-terminal region - Images



WB Suggested Anti-CTR9 Antibody Titration: 1.0 µg/ml
Positive Control: Jurkat Whole CellCTR9 is supported by BioGPS gene expression data to be expressed in Jurkat

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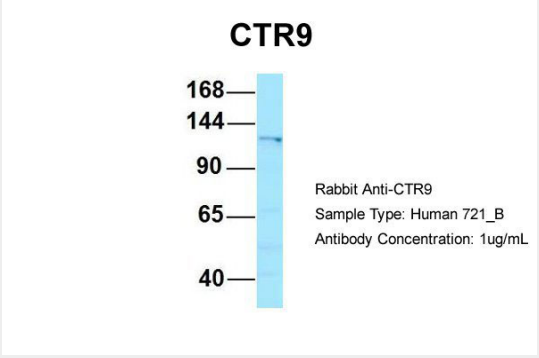

Host:Rabbit
Target Name:CTR9
Sample Tissue:Human 721_B
Antibody Dilution: 1.0µg/mlCTR9 is supported by BioGPS gene expression data to be expressed in 721_B

CTR9 antibody - C-terminal region - References

Nagase T.,et al.DNA Res. 2:167-174(1995).
Ohara O.,et al.Submitted (AUG-1995) to the EMBL/GenBank/DDBJ databases.
Mural R.J.,et al.Submitted (SEP-2005) to the EMBL/GenBank/DDBJ databases.
Zhu B.,et al.Genes Dev. 19:1668-1673(2005).
Zhu B.,et al.Mol. Cell 20:601-611(2005).