

Product datasheet

Anti-WSTF antibody [EPR1703] ab109439

KO VALIDATED Recombinant RabMAB[®]

[3 Images](#)

Overview

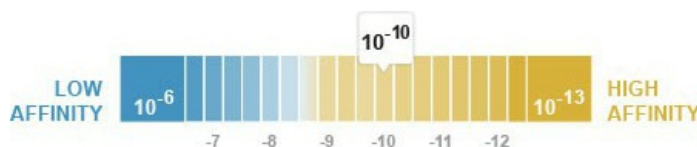
Product name	Anti-WSTF antibody [EPR1703]
Description	Rabbit monoclonal [EPR1703] to WSTF
Host species	Rabbit
Tested applications	Suitable for: WB, Flow Cyt Unsuitable for: ICC, IHC-P or IP
Species reactivity	Reacts with: Mouse, Rat, Human
Immunogen	Synthetic peptide within Human WSTF aa 1-100. The exact sequence is proprietary.
Positive control	293T, HeLa, HT-1080, PC-12, and SH-SY5Y cell lysates.
General notes	

Our RabMAB[®] technology is a patented hybridoma-based technology for making rabbit monoclonal antibodies. For details on our patents, please refer to [RabMab[®] patents](#)

This product is a recombinant rabbit monoclonal antibody.

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 = 1.23 x 10 ⁻¹⁰ M



[Learn more about K_D](#)

Storage buffer	PBS 49%, Sodium azide 0.01%, Glycerol 50%, BSA 0.05%
Purity	Tissue culture supernatant
Clonality	Monoclonal
Clone number	EPR1703

Isotype

IgG

Applications

Our [Abpromise guarantee](#) covers the use of **ab109439** 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 - 1/10000. Detects a band of approximately 185 kDa (predicted molecular weight: 171 kDa).
Flow Cyt		1/10 - 1/100. ab172730 - Rabbit monoclonal IgG, is suitable for use as an isotype control with this antibody.

Application notes

Is unsuitable for ICC, IHC-P or IP.

Target

Function

Atypical tyrosine-protein kinase that plays a central role in chromatin remodeling and acts as a transcription regulator. Involved in DNA damage response by phosphorylating 'Tyr-142' of histone H2AX (H2AXY142ph). H2AXY142ph plays a central role in DNA repair and acts as a mark that distinguishes between apoptotic and repair responses to genotoxic stress. Essential component of the WICH complex, a chromatin remodeling complex that mobilizes nucleosomes and reconfigures irregular chromatin to a regular nucleosomal array structure. The WICH complex regulates the transcription of various genes, has a role in RNA polymerase I and RNA polymerase III transcription, mediates the histone H2AX phosphorylation at 'Tyr-142', and is involved in the maintenance of chromatin structures during DNA replication processes. In the complex, it mediates the recruitment of the WICH complex to replication foci during DNA replication. Also involved in vitamin D-coupled transcription regulation via its association with the WINAC complex, a chromatin-remodeling complex recruited by vitamin D receptor (VDR), which is required for the ligand-bound VDR-mediated transrepression of the CYP27B1 gene. In the WINAC complex, plays an essential role by targeting the complex to acetylated histones, an essential step for VDR-promoter association.

Tissue specificity

Ubiquitously expressed with high levels of expression in heart, brain, placenta, skeletal muscle and ovary.

Involvement in disease

Note=BAZ1B is located in the Williams-Beuren syndrome (WBS) critical region. WBS results from a hemizygous deletion of several genes on chromosome 7q11.23, thought to arise as a consequence of unequal crossing over between highly homologous low-copy repeat sequences flanking the deleted region. Haploinsufficiency of BAZ1B may be the cause of certain cardiovascular and musculo-skeletal abnormalities observed in the disease.

Sequence similarities

Belongs to the WAL family. BAZ1B subfamily.
Contains 1 bromo domain.
Contains 1 DDT domain.
Contains 1 PHD-type zinc finger.
Contains 1 WAC domain.

Developmental stage

Expressed at equal levels in 19-23 weeks old fetal tissues.

Domain

The N-terminal part (1-345), including the WAC domain and the C motif, mediates the tyrosine-

protein kinase activity.

The bromo domain mediates the specific interaction with acetylated histones.

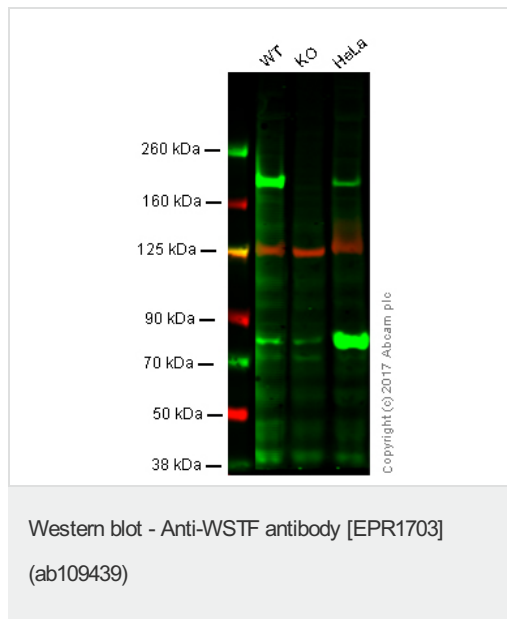
Post-translational modifications

Phosphorylated upon DNA damage, probably by ATM or ATR.

Cellular localization

Nucleus. Accumulates in pericentromeric heterochromatin during replication. Targeted to replication foci throughout S phase via its association with PCNA.

Images



Lane 1: Wild type HAP1 whole cell lysate (40 μ g)

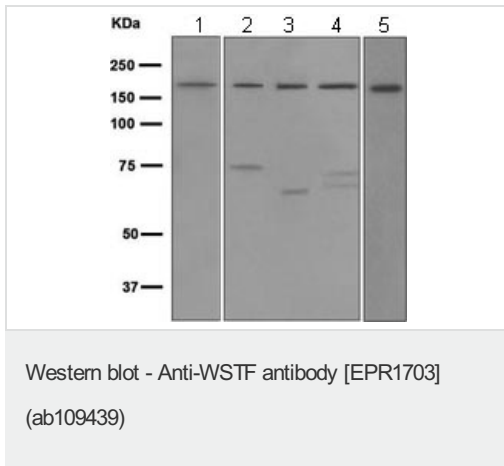
Lane 2: BAZ1B knockout HAP1 whole cell lysate (40 μ g)

Lane 3: HeLa whole cell lysate (40 μ g)

Lanes 1 - 3: Merged signal (red and green).

Green - ab109439 observed at 171 kDa. Red - loading control, ab18058, observed at 130 kDa.

Ab109439 was shown to recognize BAZ1B in wild-type cells along with additional cross-reactive bands as signal was lost in BAZ1B knockout samples. Wild-type and BAZ1B knockout samples were subjected to SDS-PAGE. Ab109439 and ab18058 (Mouse anti Vinculin loading control) were incubated overnight at 4°C at 1000 dilution and 1/10000 dilution respectively. Blots were developed with Goat anti-Rabbit IgG H&L (IRDye® 800CW) preabsorbed ab216773 and Goat anti-Mouse IgG H&L (IRDye® 680RD) preabsorbed ab216776 secondary antibodies at 1/10000 dilution for 1 hour at room temperature before imaging.



All lanes : Anti-WSTF antibody [EPR1703] (ab109439) at 1/1000 dilution

Lane 1 : 293T cell lysates

Lane 2 : HeLa cell lysates

Lane 3 : HT-1080 cell lysates

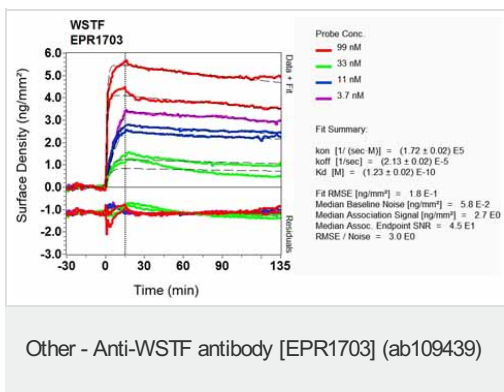
Lane 4 : PC-12 cell lysates

Lane 5 : SH-SY5Y cell lysates

Lysates/proteins at 10 µg per lane.

Predicted band size: 171 kDa

Observed band size: 185 kDa



Equilibrium dissociation constant (K_D)

Learn more about K_D

[Click here to learn more about \$K_D\$](#)

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