

Datasheet for ABIN7599753 anti-KIAA1045 antibody (AA 113-397)



Overview

Quantity:	100 μg
Target:	KIAA1045
Binding Specificity:	AA 113-397
Reactivity:	Human, Rat
Host:	Rabbit
Clonality:	Polyclonal
Conjugate:	This KIAA1045 antibody is un-conjugated
Application:	Western Blotting (WB), ELISA, Flow Cytometry (FACS)

Product Details

Purpose:	Anti-PHF24 Antibody Picoband®
Immunogen:	E.coli-derived human PHF24 recombinant protein (Position: D113-K397). Human PHF24 shares 89.5% amino acid (aa) sequence identity with mouse PHF24.
Characteristics:	Anti-PHF24 Antibody Picoband® (ABIN7599753). Tested in WB, Flow Cytometry, ELISA applications. This antibody reacts with Human, Rat. The brand Picoband indicates this is a premium antibody that guarantees superior quality, high affinity, and strong signals with minimal background in Western blot applications. Only our best-performing antibodies are designated as Picoband, ensuring unmatched performance.
Purification:	Immunogen affinity purified.

Target Details

Target:	KIAA1045
Alternative Name:	PHF24 (KIAA1045 Products)
Background:	Zinc-finger proteins contain DNA-binding domains and have a wide variety of functions, most o
	which encompass some form of transcriptional activation or repression. PHF24 (PHD finger
	protein 24), also known as KIAA1045, is a 400 amino acid protein containing one PHD-type zinc
	finger motif, suggesting involvement in transcriptional regulation events. Conserved in
	chimpanzee, Rhesus monkey, canine, bovine, mouse, rat, chicken, zebrafish and frog, the gene
	encoding PHF24 is located on human chromosome 9. Chromosome 9 houses over 900 genes
	and comprises nearly 4 % of the human genome. Considered to play a role in gender
	determination, deletion of the distal portion of chromosome 9p can lead to development of
	male to female sex reversal, the phenotype of a female with a male XY genotype.
Molecular Weight:	45 kDa
Gene ID:	23349
Application Details	
Application Notes:	Western blot, 0.25-0.5 μg/mL, Human, Rat
	Flow Cytometry (Fixed), 1-3 µg/1x10 ⁶ cells, Human
	Flow Cytometry (Fixed), 1-3 μg/1x10 ⁶ cells, Human ELISA, 0.1-0.5 μg/mL, -
	ELISA, 0.1-0.5 μg/mL, -
	ELISA, 0.1-0.5 μg/mL, - 1. Gross, M. B. Personal Communication. Baltimore, Md. 6/23/2022. 2. Kikuno, R., Nagase, T.,
	ELISA, 0.1-0.5 μg/mL, - 1. Gross, M. B. Personal Communication. Baltimore, Md. 6/23/2022. 2. Kikuno, R., Nagase, T., Ishikawa, K., Hirosawa, M., Miyajima, N., Tanaka, A., Kotani, H., Nomura, N., Ohara, O. Prediction
	ELISA, 0.1-0.5 μg/mL, - 1. Gross, M. B. Personal Communication. Baltimore, Md. 6/23/2022. 2. Kikuno, R., Nagase, T., Ishikawa, K., Hirosawa, M., Miyajima, N., Tanaka, A., Kotani, H., Nomura, N., Ohara, O. Prediction of the coding sequences of unidentified human genes. XIV. The complete sequences of 100
	ELISA, 0.1-0.5 μg/mL, - 1. Gross, M. B. Personal Communication. Baltimore, Md. 6/23/2022. 2. Kikuno, R., Nagase, T., Ishikawa, K., Hirosawa, M., Miyajima, N., Tanaka, A., Kotani, H., Nomura, N., Ohara, O. Prediction of the coding sequences of unidentified human genes. XIV. The complete sequences of 100 new cDNA clones from brain which code for large proteins in vitro. DNA Res. 6: 197-205, 1999.
Restrictions:	ELISA, 0.1-0.5 μg/mL, - 1. Gross, M. B. Personal Communication. Baltimore, Md. 6/23/2022. 2. Kikuno, R., Nagase, T., Ishikawa, K., Hirosawa, M., Miyajima, N., Tanaka, A., Kotani, H., Nomura, N., Ohara, O. Prediction of the coding sequences of unidentified human genes. XIV. The complete sequences of 100 new cDNA clones from brain which code for large proteins in vitro. DNA Res. 6: 197-205, 1999. 3. Miyamoto, K., Yamashita, A., Saito, K. Solution structure of the PHD finger from the human
Restrictions: Handling	ELISA, 0.1-0.5 µg/mL, - 1. Gross, M. B. Personal Communication. Baltimore, Md. 6/23/2022. 2. Kikuno, R., Nagase, T., Ishikawa, K., Hirosawa, M., Miyajima, N., Tanaka, A., Kotani, H., Nomura, N., Ohara, O. Prediction of the coding sequences of unidentified human genes. XIV. The complete sequences of 100 new cDNA clones from brain which code for large proteins in vitro. DNA Res. 6: 197-205, 1999. 3. Miyamoto, K., Yamashita, A., Saito, K. Solution structure of the PHD finger from the human KIAA1045 protein. Protein Sci. 27: 987-992, 2018.
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Format:	Lyophilized
Reconstitution:	Adding 0.2 mL of distilled water will yield a concentration of 500 $\mu g/mL$.
Concentration:	500 μg/mL
Buffer:	Each vial contains 4 mg Trehalose, 0.9 mg NaCl, 0.2 mg Na2HPO4.
Storage:	4 °C,-20 °C

Handling

Storage Comment:

At -20°C for one year from date of receipt. After reconstitution, at 4°C for one month.

It can also be aliquotted and stored frozen at -20°C for six months. Avoid repeated freezing and thawing.