

Resource Summary Report

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Rabbit TrueBlot: Anti-Rabbit IgG HRP - 18-8816-33

RRID:AB_2610848

Type: Antibody

Proper Citation

(Rockland Cat# 18-8816-33, RRID:AB_2610848)

Antibody Information

URL: http://antibodyregistry.org/AB_2610848

Proper Citation: (Rockland Cat# 18-8816-33, RRID:AB_2610848)

Target Antigen: Rabbit TrueBlot: Anti-Rabbit IgG HRP

Host Organism: mouse

Clonality: monoclonal

Comments: Western Blot, Rabbit IgG TrueBlot is provided as 1000X solution

Antibody Name: Rabbit TrueBlot: Anti-Rabbit IgG HRP - 18-8816-33

Description: This monoclonal targets Rabbit TrueBlot: Anti-Rabbit IgG HRP

Target Organism: rabbit

Clone ID: eB182

Antibody ID: AB_2610848

Vendor: Rockland

Catalog Number: 18-8816-33

Record Creation Time: 20231110T035003+0000

Record Last Update: 20240725T002726+0000

Ratings and Alerts

No rating or validation information has been found for Rabbit TrueBlot: Anti-Rabbit IgG HRP - 18-8816-33.

No alerts have been found for Rabbit TrueBlot: Anti-Rabbit IgG HRP - 18-8816-33.

Data and Source Information

Source: [Antibody Registry](#)

Usage and Citation Metrics

We found 31 mentions in open access literature.

Listed below are recent publications. The full list is available at [FDI Lab - SciCrunch.org](#).

Surana S, et al. (2024) The tyrosine phosphatases LAR and PTPR γ act as receptors of the nidogen-tetanus toxin complex. The EMBO journal, 43(16), 3358.

Bossaert M, et al. (2024) Identification of the main barriers to Ku accumulation in chromatin. Cell reports, 43(8), 114538.

Matsui S, et al. (2024) Pioneer and PRDM transcription factors coordinate bivalent epigenetic states to safeguard cell fate. Molecular cell, 84(3), 476.

Deka A, et al. (2024) Non-canonical NF- κ B signaling limits the tolerogenic β -catenin-Raldh2 axis in gut dendritic cells to exacerbate intestinal pathologies. The EMBO journal, 43(18), 3895.

Scholtes C, et al. (2024) Identification of a chromatin-bound ERR α interactome network in mouse liver. Molecular metabolism, 83, 101925.

Fang Z, et al. (2023) Aurora A polyubiquitinates the BRCA1-interacting protein OLA1 to promote centrosome maturation. Cell reports, 42(8), 112850.

Boehm D, et al. (2023) The lysine methyltransferase SMYD5 amplifies HIV-1 transcription and is post-transcriptionally upregulated by Tat and USP11. Cell reports, 42(3), 112234.

Refaat AM, et al. (2023) HNRNPU facilitates antibody class-switch recombination through C-NHEJ promotion and R-loop suppression. Cell reports, 42(3), 112284.

He H, et al. (2023) PRDM3/16 Regulate Chromatin Accessibility Required for NKX2-1 Mediated Alveolar Epithelial Differentiation and Function. bioRxiv : the preprint server for biology.

Aryan F, et al. (2023) Nucleolus activity-dependent recruitment and biomolecular condensation by pH sensing. Molecular cell, 83(23), 4413.

Janneh AH, et al. (2022) Crosstalk between pro-survival sphingolipid metabolism and complement signaling induces inflammasome-mediated tumor metastasis. *Cell reports*, 41(10), 111742.

Liang G, et al. (2022) CTNNB1 restricts HIV-1 replication by suppressing viral DNA integration into the cell genome. *Cell reports*, 38(11), 110533.

Liu X, et al. (2022) The PAF1 complex promotes 3' processing of pervasive transcripts. *Cell reports*, 38(11), 110519.

Blue SM, et al. (2022) Transcriptome-wide identification of RNA-binding protein binding sites using seCLIP-seq. *Nature protocols*, 17(5), 1223.

Alerasool N, et al. (2022) Identification and functional characterization of transcriptional activators in human cells. *Molecular cell*, 82(3), 677.

Kelenis DP, et al. (2022) Inhibition of Karyopherin β 1-Mediated Nuclear Import Disrupts Oncogenic Lineage-Defining Transcription Factor Activity in Small Cell Lung Cancer. *Cancer research*, 82(17), 3058.

Kawashima T, et al. (2021) Phosphorylation of Collapsin Response Mediator Protein 1 (CRMP1) at Tyrosine 504 residue regulates Semaphorin 3A-induced cortical dendritic growth. *Journal of neurochemistry*, 157(4), 1207.

Corley M, et al. (2021) fSHAPE, fSHAPE-eCLIP, and SHAPE-eCLIP probe transcript regions that interact with specific proteins. *STAR protocols*, 2(3), 100762.

Piette BL, et al. (2021) Comprehensive interactome profiling of the human Hsp70 network highlights functional differentiation of J domains. *Molecular cell*, 81(12), 2549.

Reichermeier KM, et al. (2020) PIKES Analysis Reveals Response to Degradation and Key Regulatory Mechanisms of the CRL4 Network. *Molecular cell*, 77(5), 1092.