

Resource Summary Report

Generated by [RRID](#) on Jul 30, 2026

Rabbit anti-RNF2 Antibody, Affinity Purified

RRID:AB_10632773

Type: Antibody

Proper Citation

Bethyl Cat# A302-869A, RRID:AB_10632773

Antibody Information

URL: http://antibodyregistry.org/AB_10632773

Proper Citation: Bethyl Cat# A302-869A, RRID:AB_10632773

Target Antigen: RNF2

Host Organism: rabbit

Clonality: polyclonal

Comments: Applications: WB, IP, IHC, ChIP-Seq

Antibody Name: Rabbit anti-RNF2 Antibody, Affinity Purified

Description: This polyclonal targets RNF2

Target Organism: mouse, human

Antibody ID: AB_10632773

Vendor: Bethyl

Catalog Number: A302-869A

Alternative Catalog Numbers: A302-869A-M, A302-869A-T

Record Creation Time: 20250820T024333+0000

Record Last Update: 20250820T142110+0000

Ratings and Alerts

- ENCODE PROJECT External validation for lot: 1 is available under ENCODE ID: ENCAB790JUW - ENCODE
<https://www.encodeproject.org/antibodies/ENCAB790JUW>

Warning: Discontinued at Thermo Fisher Scientific
Applications: WB, IP, IHC, ChIP-Seq

Data and Source Information

Source: [Antibody Registry](#)

Usage and Citation Metrics

We found 8 mentions in open access literature.

Listed below are recent publications. The full list is available at [RRID](#) .

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

Matsui S, et al. (2024) Protocol for establishing inducible CRISPRd system for blocking transcription factor-binding sites in human pluripotent stem cells. *STAR protocols*, 5(3), 103233.

Liang Q, et al. (2022) Essential role of MESP1-RING1A complex in cardiac differentiation. *Developmental cell*, 57(22), 2533.

Liu S, et al. (2021) NRF1 association with AUTS2-Polycomb mediates specific gene activation in the brain. *Molecular cell*, 81(22), 4663.

Miyamoto R, et al. (2020) Activation of CpG-Rich Promoters Mediated by MLL Drives MOZ-Rearranged Leukemia. *Cell reports*, 32(13), 108200.

Wang Z, et al. (2018) A Non-canonical BCOR-PRC1.1 Complex Represses Differentiation Programs in Human ESCs. *Cell stem cell*, 22(2), 235.

Lyu X, et al. (2018) Architectural Proteins and Pluripotency Factors Cooperate to Orchestrate the Transcriptional Response of hESCs to Temperature Stress. *Molecular cell*, 71(6), 940.

Kundu S, et al. (2017) Polycomb Repressive Complex 1 Generates Discrete Compacted Domains that Change during Differentiation. *Molecular cell*, 65(3), 432.