

Resource Summary Report

Generated by [RRID](#) on Sep 11, 2026

[RbAp48 antibody \[EPR3411\]](#)

RRID:AB_1603754

Type: Antibody

Proper Citation

Abcam Cat# ab79416, RRID:AB_1603754

Antibody Information

URL: http://antibodyregistry.org/AB_1603754

Proper Citation: Abcam Cat# ab79416, RRID:AB_1603754

Target Antigen: RbAp48 antibody [EPR3411]

Host Organism: rabbit

Clonality: monoclonal

Comments: validation status unknown, seller recommendations provided in 2012: Flow Cyt, ICC/IF, IHC-P, IP, WB; Immunohistochemistry - fixed; Flow Cytometry; Immunocytochemistry; Immunoprecipitation; Immunofluorescence; Immunohistochemistry; Western Blot

Antibody Name: RbAp48 antibody [EPR3411]

Description: This monoclonal targets RbAp48 antibody [EPR3411]

Target Organism: rat, mouse, human

Antibody ID: AB_1603754

Vendor: Abcam

Catalog Number: ab79416

Record Creation Time: 20231110T073516+0000

Record Last Update: 20260829T012011+0000

Ratings and Alerts

No rating or validation information has been found for RbAp48 antibody [EPR3411].

No alerts have been found for RbAp48 antibody [EPR3411].

Data and Source Information

Source: [Antibody Registry](#)

Usage and Citation Metrics

We found 5 mentions in open access literature.

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

Liu N, et al. (2023) Histone H3 lysine 27 crotonylation mediates gene transcriptional repression in chromatin. *Molecular cell*, 83(13), 2206.

Cao X, et al. (2021) Circular RNA hsa_circ_0102231 sponges miR-145 to promote non-small cell lung cancer cell proliferation by up-regulating the expression of RBBP4. *Journal of biochemistry*, 169(1), 65.

Miao X, et al. (2020) Loss of RBBP4 results in defective inner cell mass, severe apoptosis, hyperacetylated histones and preimplantation lethality in mice†. *Biology of reproduction*, 103(1), 13.

Gomes AP, et al. (2019) Dynamic Incorporation of Histone H3 Variants into Chromatin Is Essential for Acquisition of Aggressive Traits and Metastatic Colonization. *Cancer cell*, 36(4), 402.

Bornelöv S, et al. (2018) The Nucleosome Remodeling and Deacetylation Complex Modulates Chromatin Structure at Sites of Active Transcription to Fine-Tune Gene Expression. *Molecular cell*, 71(1), 56.