Resource Summary Report

Generated by RRID on Aug 9, 2026

SMARCC2/BAF170 (D8O9V) Rabbit mAb

RRID:AB_2798017
Type: Antibody

Proper Citation

Cell Signaling Technology Cat# 12760, RRID:AB_2798017

Antibody Information

URL: http://antibodyregistry.org/AB_2798017

Proper Citation: Cell Signaling Technology Cat# 12760, RRID:AB_2798017

Target Antigen: SMARCC2

Host Organism: rabbit

Clonality: monoclonal

Comments: Applications: W, IP, ChIP, ChIP-seq

Antibody Name: SMARCC2/BAF170 (D8O9V) Rabbit mAb

Description: This monoclonal targets SMARCC2

Target Organism: h, m, r, mk

Clone ID: Clone D8O9V

Antibody ID: AB_2798017

Vendor: Cell Signaling Technology

Catalog Number: 12760

Record Creation Time: 20250820T054447+0000

Record Last Update: 20250820T222140+0000

Ratings and Alerts

No rating or validation information has been found for SMARCC2/BAF170 (D8O9V) Rabbit mAb.

No alerts have been found for SMARCC2/BAF170 (D8O9V) Rabbit mAb.

Data and Source Information

Source: Antibody Registry

Usage and Citation Metrics

We found 7 mentions in open access literature.

Listed below are recent publications. The full list is available at RRID .

Ma X, et al. (2026) IAP-based biodegraders can convert necroptosis to apoptosis and eliminate cancer driving protein complexes. Cell chemical biology, 33(4), 553.

Duplaquet L, et al. (2024) Mammalian SWI/SNF complex activity regulates POU2F3 and constitutes a targetable dependency in small cell lung cancer. Cancer cell, 42(8), 1352.

Zhang C, et al. (2024) Methionine secreted by tumor-associated pericytes supports cancer stem cells in clear cell renal carcinoma. Cell metabolism, 36(4), 778.

Liu W, et al. (2023) RNF138 inhibits late inflammatory gene transcription through degradation of SMARCC1 of the SWI/SNF complex. Cell reports, 42(2), 112097.

Pintacuda G, et al. (2023) Protein interaction studies in human induced neurons indicate convergent biology underlying autism spectrum disorders. Cell genomics, 3(3), 100250.

Patil A, et al. (2023) A disordered region controls cBAF activity via condensation and partner recruitment. Cell, 186(22), 4936.

Simeoni F, et al. (2021) Enhancer recruitment of transcription repressors RUNX1 and TLE3 by mis-expressed FOXC1 blocks differentiation in acute myeloid leukemia. Cell reports, 36(12), 109725.