

# Resource Summary Report

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

## Rabbit Anti-SRC-3 Monoclonal Antibody, Unconjugated, Clone 5E11

RRID:AB\_823642

Type: Antibody

### Proper Citation

Cell Signaling Technology Cat# 2126, RRID:AB\_823642

### Antibody Information

**URL:** [http://antibodyregistry.org/AB\\_823642](http://antibodyregistry.org/AB_823642)

**Proper Citation:** Cell Signaling Technology Cat# 2126, RRID:AB\_823642

**Target Antigen:** SRC-3

**Host Organism:** rabbit

**Clonality:** monoclonal

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

**Antibody Name:** Rabbit Anti-SRC-3 Monoclonal Antibody, Unconjugated, Clone 5E11

**Description:** This monoclonal targets SRC-3

**Target Organism:** monkey, rat, simian, mouse, human

**Clone ID:** Clone 5E11

**Antibody ID:** AB\_823642

**Vendor:** Cell Signaling Technology

**Catalog Number:** 2126

**Record Creation Time:** 20250820T014156+0000

**Record Last Update:** 20250820T113731+0000

### Ratings and Alerts

No rating or validation information has been found for Rabbit Anti-SRC-3 Monoclonal Antibody, Unconjugated, Clone 5E11.

No alerts have been found for Rabbit Anti-SRC-3 Monoclonal Antibody, Unconjugated, Clone 5E11.

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## Data and Source Information

**Source:** [Antibody Registry](#)

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## Usage and Citation Metrics

We found 5 mentions in open access literature.

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

Takahashi M, et al. (2025) PPM1D is directly degraded by proteasomes in a ubiquitination-independent manner through its carboxyl-terminal region. Journal of biomedical science, 32(1), 88.

Han SJ, et al. (2023) Steroid Receptor Coactivator-3 is a Key Modulator of Regulatory T Cell-Mediated Tumor Evasion. bioRxiv : the preprint server for biology.

Meng H, et al. (2022) Defining the mammalian coactivation of hepatic 12-h clock and lipid metabolism. Cell reports, 38(10), 110491.

Sharif GM, et al. (2021) An AIB1 Isoform Alters Enhancer Access and Enables Progression of Early-Stage Triple-Negative Breast Cancer. Cancer research, 81(16), 4230.

Vilariño-Güell C, et al. (2019) Exome sequencing in multiple sclerosis families identifies 12 candidate genes and nominates biological pathways for the genesis of disease. PLoS genetics, 15(6), e1008180.