

# Resource Summary Report

Generated by [FDI Lab - SciCrunch.org](https://www.fdi-lab.com) on Apr 13, 2025

## Myc-Tag (9B11) Mouse mAb (Alexa Fluor 488 Conjugate)

RRID:AB\_2151849

Type: Antibody

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### Proper Citation

(Cell Signaling Technology Cat# 2279, RRID:AB\_2151849)

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### Antibody Information

**URL:** [http://antibodyregistry.org/AB\\_2151849](http://antibodyregistry.org/AB_2151849)

**Proper Citation:** (Cell Signaling Technology Cat# 2279, RRID:AB\_2151849)

**Target Antigen:** Myc-Tag (9B11) Mouse mAb (Alexa Fluor 488 Conjugate)

**Host Organism:** mouse

**Clonality:** monoclonal

**Comments:** Applications: IF-IC, F. Consolidation on 10/2018: AB\_10693383, AB\_2151849.

**Antibody Name:** Myc-Tag (9B11) Mouse mAb (Alexa Fluor 488 Conjugate)

**Description:** This monoclonal targets Myc-Tag (9B11) Mouse mAb (Alexa Fluor 488 Conjugate)

**Target Organism:** all

**Antibody ID:** AB\_2151849

**Vendor:** Cell Signaling Technology

**Catalog Number:** 2279

**Record Creation Time:** 20241017T004235+0000

**Record Last Update:** 20241017T023523+0000

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## Ratings and Alerts

No rating or validation information has been found for Myc-Tag (9B11) Mouse mAb (Alexa Fluor 488 Conjugate).

No alerts have been found for Myc-Tag (9B11) Mouse mAb (Alexa Fluor 488 Conjugate).

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 4 mentions in open access literature.

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

Kudo T, et al. (2022) A multiplexed epitope barcoding strategy that enables dynamic cellular phenotypic screens. *Cell systems*, 13(5), 376.

Glassman CR, et al. (2021) Structural basis for IL-12 and IL-23 receptor sharing reveals a gateway for shaping actions on T versus NK cells. *Cell*, 184(4), 983.

Cai R, et al. (2021) Ion permeation controlled by hydrophobic residues and proton binding in the proton-activated chloride channel. *iScience*, 24(12), 103395.

Mendoza JL, et al. (2020) Interrogating the recognition landscape of a conserved HIV-specific TCR reveals distinct bacterial peptide cross-reactivity. *eLife*, 9.