

Resource Summary Report

Generated by [RRID](#) on Sep 6, 2026

CD8a

RRID:AB_396780

Type: Antibody

Proper Citation

BD Biosciences Cat# 557668, RRID:AB_396780

Antibody Information

URL: http://antibodyregistry.org/AB_396780

Proper Citation: BD Biosciences Cat# 557668, RRID:AB_396780

Target Antigen: CD8a

Host Organism: rat

Clonality: monoclonal

Comments: Applications: Immunofluorescence, Flow cytometry

Antibody Name: CD8a

Description: This monoclonal targets CD8a

Target Organism: mouse

Antibody ID: AB_396780

Vendor: BD Biosciences

Catalog Number: 557668

Record Creation Time: 20231110T081136+0000

Record Last Update: 20260831T222828+0000

Ratings and Alerts

No rating or validation information has been found for CD8a.

No alerts have been found for CD8a.

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](#) .

Libri AB, et al. (2025) Senataxin promotes recombination fidelity during antigen receptor gene diversification. Science signaling, 18(908), eadv8801.

Archer D, et al. (2025) A secondary metabolite of Limosilactobacillusreuteri R2lc drives strain-specific pathology in a spontaneous mouse model of multiple sclerosis. Cell reports, 44(3), 115321.

Archer D, et al. (2023) The importance of the timing of microbial signals for perinatal immune system development. Microbiome research reports, 2(2), 11.

VanDyke D, et al. (2022) Engineered human cytokine/antibody fusion proteins expand regulatory T??cells and confer autoimmune disease protection. Cell reports, 41(3), 111478.

Xia K, et al. (2022) AAV-mediated gene therapy produces fertile offspring in the Lhcgr-deficient mouse model of Leydig cell failure. Cell reports. Medicine, 3(11), 100792.

Ahrends T, et al. (2021) Enteric pathogens induce tissue tolerance and prevent neuronal loss from subsequent infections. Cell, 184(23), 5715.

Bianchi JJ, et al. (2019) Breakage-Fusion-Bridge Events Trigger Complex Genome Rearrangements and Amplifications in Developmentally Arrested T Cell Lymphomas. Cell reports, 27(10), 2847.