Resource Summary Report

Generated by FDI Lab - SciCrunch.org on May 23, 2025

TotalSeq(TM)-A0304 anti-mouse Hashtag 4

RRID:AB_2750035
Type: Antibody

Proper Citation

(BioLegend Cat# 155807, RRID:AB_2750035)

Antibody Information

URL: http://antibodyregistry.org/AB_2750035

Proper Citation: (BioLegend Cat# 155807, RRID:AB_2750035)

Target Antigen: H-2/CD45

Host Organism: rat

Clonality: monoclonal

Comments: Applications: PG

Antibody Name: TotalSeq(TM)-A0304 anti-mouse Hashtag 4

Description: This monoclonal targets H-2/CD45

Target Organism: mouse

Clone ID: Clone M1/42; 30-F11

Antibody ID: AB_2750035

Vendor: BioLegend

Catalog Number: 155807

Record Creation Time: 20231110T033401+0000

Record Last Update: 20240725T020543+0000

Ratings and Alerts

No rating or validation information has been found for TotalSeq(TM)-A0304 anti-mouse Hashtag 4.

No alerts have been found for TotalSeq(TM)-A0304 anti-mouse Hashtag 4.

Data and Source Information

Source: Antibody Registry

Usage and Citation Metrics

We found 17 mentions in open access literature.

Listed below are recent publications. The full list is available at FDI Lab - SciCrunch.org.

Sacirbegovic F, et al. (2023) Graft-versus-host disease is locally maintained in target tissues by resident progenitor-like T cells. Immunity, 56(2), 369.

Grootveld AK, et al. (2023) Apoptotic cell fragments locally activate tingible body macrophages in the germinal center. Cell, 186(6), 1144.

Ugur M, et al. (2023) Lymph node medulla regulates the spatiotemporal unfolding of resident dendritic cell networks. Immunity, 56(8), 1778.

Konturek-Ciesla A, et al. (2023) Temporal multimodal single-cell profiling of native hematopoiesis illuminates altered differentiation trajectories with age. Cell reports, 42(4), 112304.

Hou P, et al. (2023) The ?-secretase substrate proteome and its role in cell signaling regulation. Molecular cell, 83(22), 4106.

Kotov DI, et al. (2023) Early cellular mechanisms of type I interferon-driven susceptibility to tuberculosis. Cell, 186(25), 5536.

Tessaro FHG, et al. (2022) Single-cell RNA-seq of a soft-tissue sarcoma model reveals the critical role of tumor-expressed MIF in shaping macrophage heterogeneity. Cell reports, 39(12), 110977.

Širvinskas D, et al. (2022) Single-cell atlas of the aging mouse colon. iScience, 25(5), 104202.

Ben-Moshe S, et al. (2022) The spatiotemporal program of zonal liver regeneration following acute injury. Cell stem cell, 29(6), 973.

Milon B, et al. (2021) A cell-type-specific atlas of the inner ear transcriptional response to acoustic trauma. Cell reports, 36(13), 109758.

Fast EM, et al. (2021) External signals regulate continuous transcriptional states in hematopoietic stem cells. eLife, 10.

Pankaew S, et al. (2021) Multiplexed single-cell RNA-sequencing of mouse thymic and splenic samples. STAR protocols, 3(1), 101041.

Guldner IH, et al. (2021) Isolation of mouse brain-infiltrating leukocytes for single cell profiling of epitopes and transcriptomes. STAR protocols, 2(2), 100537.

Nozais M, et al. (2021) MYC deficiency impairs the development of effector/memory T lymphocytes. iScience, 24(7), 102761.

Guldner IH, et al. (2020) CNS-Native Myeloid Cells Drive Immune Suppression in the Brain Metastatic Niche through Cxcl10. Cell, 183(5), 1234.

Golomb SM, et al. (2020) Multi-modal Single-Cell Analysis Reveals Brain Immune Landscape Plasticity during Aging and Gut Microbiota Dysbiosis. Cell reports, 33(9), 108438.

Soliman H, et al. (2020) Pathogenic Potential of Hic1-Expressing Cardiac Stromal Progenitors. Cell stem cell, 26(2), 205.