

# Resource Summary Report

Generated by [dkNET](#) on Sep 3, 2026

## TotalSeq(TM)-A0253 anti-human Hashtag 3

RRID:AB\_2750017

Type: Antibody

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### Proper Citation

BioLegend Cat# 394605, RRID:AB\_2750017

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### Antibody Information

**URL:** [http://antibodyregistry.org/AB\\_2750017](http://antibodyregistry.org/AB_2750017)

**Proper Citation:** BioLegend Cat# 394605, RRID:AB\_2750017

**Target Antigen:** CD298/beta2-microglobulin

**Host Organism:** mouse

**Clonality:** monoclonal

**Comments:** Applications: PG

**Antibody Name:** TotalSeq(TM)-A0253 anti-human Hashtag 3

**Description:** This monoclonal targets CD298/beta2-microglobulin

**Target Organism:** human

**Clone ID:** Clone LNH-94; 2M2

**Antibody ID:** AB\_2750017

**Vendor:** BioLegend

**Catalog Number:** 394605

**Record Creation Time:** 20231110T033401+0000

**Record Last Update:** 20260829T183515+0000

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### Ratings and Alerts

No rating or validation information has been found for TotalSeq(TM)-A0253 anti-human Hashtag 3.

No alerts have been found for TotalSeq(TM)-A0253 anti-human Hashtag 3.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 24 mentions in open access literature.

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

Rigby RE, et al. (2026) Single-cell analysis of signalling and transcriptional responses to type I interferons. EMBO reports, 27(9), 2491.

Kremneva E, et al. (2026) Precision tumor-on-chip for personalized assessment of drug efficacy and immune cell delivery. Cell reports methods, 101475.

Wang H, et al. (2026) An immunocompetent bone marrow-on-a-chip model for studying human hematological malignancies and preclinical therapeutic screening. Nature protocols.

Chen T, et al. (2026) Multi-modal single-cell sequencing reveals network transition in circulating monocytes that aligns with faster recovery in patients with trauma and favours a response to M-CSF. EBioMedicine, 130, 106368.

Lombardi O, et al. (2025) Conserved patterns of transcriptional dysregulation, heterogeneity, and cell states in clear cell kidney cancer. Cell reports, 44(1), 115169.

Caron DP, et al. (2025) Multimodal hierarchical classification of CITE-seq data delineates immune cell states across lineages and tissues. Cell reports methods, 5(1), 100938.

Collora JA, et al. (2025) Medications for opioid use disorder shape immune responses during chronic HIV infection. Cell reports. Medicine, 6(6), 102159.

Veronese-Paniagua DA, et al. (2025) Coxsackievirus B infection invokes unique cell-type-specific responses in primary human pancreatic islets. Cell reports, 44(9), 116211.

Wei Y, et al. (2025) Transcription factor BACH2 shapes tissue-resident memory T cell programs to promote HIV-1 persistence. Immunity.

Jaeger-Ruckstuhl CA, et al. (2024) Signaling via a CD27-TRAF2-SHP-1 axis during naive T cell activation promotes memory-associated gene regulatory networks. Immunity, 57(2), 287.

Gan S, et al. (2024) Distinct tumor architectures and microenvironments for the initiation of breast cancer metastasis in the brain. Cancer cell, 42(10), 1693.

MacDonald L, et al. (2024) Synovial tissue myeloid dendritic cell subsets exhibit distinct tissue-niche localization and function in health and rheumatoid arthritis. Immunity, 57(12), 2843.

Wei Y, et al. (2023) Single-cell epigenetic, transcriptional, and protein profiling of latent and active HIV-1 reservoir revealed that IKZF3 promotes HIV-1 persistence. *Immunity*, 56(11), 2584.

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

Krausgruber T, et al. (2023) Single-cell and spatial transcriptomics reveal aberrant lymphoid developmental programs driving granuloma formation. *Immunity*, 56(2), 289.

Lombardi O, et al. (2022) Pan-cancer analysis of tissue and single-cell HIF-pathway activation using a conserved gene signature. *Cell reports*, 41(7), 111652.

Masle-Farquhar E, et al. (2022) STAT3 gain-of-function mutations connect leukemia with autoimmune disease by pathological NKG2Dhi CD8<sup>+</sup> T cell dysregulation and accumulation. *Immunity*, 55(12), 2386.

Krämer B, et al. (2021) Early IFN- $\gamma$  signatures and persistent dysfunction are distinguishing features of NK cells in severe COVID-19. *Immunity*, 54(11), 2650.

Yao C, et al. (2021) Cell-Type-Specific Immune Dysregulation in Severely Ill COVID-19 Patients. *Cell reports*, 34(1), 108590.

Fawkner-Corbett D, et al. (2021) Spatiotemporal analysis of human intestinal development at single-cell resolution. *Cell*, 184(3), 810.