

Resource Summary Report

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

BUV661 Mouse Anti-Human CD14

RRID:AB_2868407

Type: Antibody

Proper Citation

BD Biosciences Cat# 741684, RRID:AB_2868407

Antibody Information

URL: http://antibodyregistry.org/AB_2868407

Proper Citation: BD Biosciences Cat# 741684, RRID:AB_2868407

Target Antigen: CD14

Host Organism: mouse

Clonality: monoclonal

Comments: Applications: Flow cytometry

Antibody Name: BUV661 Mouse Anti-Human CD14

Description: This monoclonal targets CD14

Target Organism: human

Clone ID: M?P9

Antibody ID: AB_2868407

Vendor: BD Biosciences

Catalog Number: 741684

Record Creation Time: 20231110T031939+0000

Record Last Update: 20260829T191828+0000

Ratings and Alerts

No rating or validation information has been found for BUV661 Mouse Anti-Human CD14.

No alerts have been found for BUV661 Mouse Anti-Human CD14.

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 [dkNET](#) .

Schuster DJ, et al. (2026) Coordinated immune response distinguishes duration of SARS-CoV-2 viral particle shedding in humans. *iScience*, 29(4), 115277.

Conn BP, et al. (2025) Generation of T cell responses against broad KRAS hotspot neoantigens for cell therapy or TCR discovery. *Cell reports methods*, 5(5), 101049.

Mayer-Blackwell K, et al. (2023) mRNA vaccination boosts S-specific T cell memory and promotes expansion of CD45RA^{int} TEMRA-like CD8⁺ T cells in COVID-19 recovered individuals. *Cell reports. Medicine*, 4(8), 101149.

Cohen KW, et al. (2021) Longitudinal analysis shows durable and broad immune memory after SARS-CoV-2 infection with persisting antibody responses and memory B and T cells. *Cell reports. Medicine*, 2(7), 100354.

Genge PC, et al. (2021) Optimized workflow for human PBMC multiomic immunosurveillance studies. *STAR protocols*, 2(4), 100900.

Swanson E, et al. (2021) Simultaneous trimodal single-cell measurement of transcripts, epitopes, and chromatin accessibility using TEA-seq. *eLife*, 10.

Savage AK, et al. (2021) Multimodal analysis for human ex vivo studies shows extensive molecular changes from delays in blood processing. *iScience*, 24(5), 102404.