

Resource Summary Report

Generated by [NIF](#) on Jul 29, 2026

Anti-Human CD14 (LPS receptor) Monoclonal Antibody, ECD Conjugated, Clone RMO52

RRID:AB_130853

Type: Antibody

Proper Citation

Beckman Coulter Cat# IM2707U, RRID:AB_130853

Antibody Information

URL: http://antibodyregistry.org/AB_130853

Proper Citation: Beckman Coulter Cat# IM2707U, RRID:AB_130853

Target Antigen: Human CD14 (LPS receptor)

Host Organism: mouse

Clonality: monoclonal

Comments: manufacturer recommendations: Flow Cytometry; Flow Cytometry

Antibody Name: Anti-Human CD14 (LPS receptor) Monoclonal Antibody, ECD Conjugated, Clone RMO52

Description: This monoclonal targets Human CD14 (LPS receptor)

Target Organism: human

Clone ID: Clone RMO52

Antibody ID: AB_130853

Vendor: Beckman Coulter

Catalog Number: IM2707U

Record Creation Time: 20250819T235808+0000

Record Last Update: 20250820T065822+0000

Ratings and Alerts

No rating or validation information has been found for Anti-Human CD14 (LPS receptor) Monoclonal Antibody, ECD Conjugated, Clone RMO52.

No alerts have been found for Anti-Human CD14 (LPS receptor) Monoclonal Antibody, ECD Conjugated, Clone RMO52.

Data and Source Information

Source: [Antibody Registry](#)

Usage and Citation Metrics

We found 5 mentions in open access literature.

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

Moorlag SJCFM, et al. (2024) Multi-omics analysis of innate and adaptive responses to BCG vaccination reveals epigenetic cell states that predict trained immunity. *Immunity*, 57(1), 171.

Wu HL, et al. (2023) Allogeneic immunity clears latent virus following allogeneic stem cell transplantation in SIV-infected ART-suppressed macaques. *Immunity*, 56(7), 1649.

Frishberg A, et al. (2022) Mature neutrophils and a NF- κ B-to-IFN transition determine the unifying disease recovery dynamics in COVID-19. *Cell reports. Medicine*, 3(6), 100652.

Emanuel KM, et al. (2022) Deprenyl reduces inflammation during acute SIV infection. *iScience*, 25(5), 104207.

Upadhyay AA, et al. (2018) BALDR: a computational pipeline for paired heavy and light chain immunoglobulin reconstruction in single-cell RNA-seq data. *Genome medicine*, 10(1), 20.