Resource Summary Report

Generated by dkNET on Aug 16, 2026

Recombinant Anti-CD14 antibody [EPR3653]

RRID:AB_2889158
Type: Antibody

Proper Citation

Abcam Cat# ab133335, RRID:AB_2889158

Antibody Information

URL: http://antibodyregistry.org/AB_2889158

Proper Citation: Abcam Cat# ab133335, RRID:AB_2889158

Target Antigen: CD14

Host Organism: rabbit

Clonality: recombinant monoclonal

Comments: Applications: WB, IHC-P

Antibody Name: Recombinant Anti-CD14 antibody [EPR3653]

Description: This recombinant monoclonal targets CD14

Target Organism: human

Clone ID: EPR3653

Antibody ID: AB_2889158

Vendor: Abcam

Catalog Number: ab133335

Record Creation Time: 20250819T205919+0000

Record Last Update: 20250819T212837+0000

Ratings and Alerts

No rating or validation information has been found for Recombinant Anti-CD14 antibody [EPR3653].

No alerts have been found for Recombinant Anti-CD14 antibody [EPR3653].

Data and Source Information

Source: Antibody Registry

Usage and Citation Metrics

We found 9 mentions in open access literature.

Listed below are recent publications. The full list is available at dkNET .

Toh J, et al. (2024) Multi-modal analysis reveals tumor and immune features distinguishing EBV-positive and EBV-negative post-transplant lymphoproliferative disorders. Cell reports. Medicine, 5(12), 101851.

Wang S, et al. (2024) Disruption of MerTK increases the efficacy of checkpoint inhibitor by enhancing ferroptosis and immune response in hepatocellular carcinoma. Cell reports. Medicine, 5(2), 101415.

van Eijs MJM, et al. (2023) Highly multiplexed spatial analysis identifies tissue-resident memory T??cells as drivers of ulcerative and immune checkpoint inhibitor colitis. iScience, 26(10), 107891.

Li J, et al. (2023) Remodeling of the immune and stromal cell compartment by PD-1 blockade in mismatch repair-deficient colorectal cancer. Cancer cell, 41(6), 1152.

Varn FS, et al. (2022) Glioma progression is shaped by genetic evolution and microenvironment interactions. Cell, 185(12), 2184.

Jarosch S, et al. (2022) ChipCytometry for multiplexed detection of protein and mRNA markers on human FFPE tissue samples. STAR protocols, 3(2), 101374.

Krijgsman D, et al. (2022) MATISSE: An analysis protocol for combining imaging mass cytometry with fluorescence microscopy to generate single-cell data. STAR protocols, 3(1), 101034.

Jarosch S, et al. (2021) Multiplexed imaging and automated signal quantification in formalin-fixed paraffin-embedded tissues by ChipCytometry. Cell reports methods, 1(7), 100104.

Fu H, et al. (2021) The hepatocyte growth factor/c-met pathway is a key determinant of the fibrotic kidney local microenvironment. iScience, 24(10), 103112.