

# Resource Summary Report

Generated by [RRID](#) on Sep 8, 2026

## Recombinant Anti-CD14 antibody [EPR3653]

RRID:AB\_2889158

Type: Antibody

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### Proper Citation

Abcam Cat# ab133335, RRID:AB\_2889158

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### Antibody Information

**URL:** [http://antibodyregistry.org/AB\\_2889158](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:** 20231110T031709+0000

**Record Last Update:** 20260828T234719+0000

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### 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].

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 9 mentions in open access literature.

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

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.

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

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.

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.