

Resource Summary Report

Generated by [RRID](#) on Sep 7, 2026

Recombinant Anti-RAB8A antibody [EPR14873]

RRID:AB_2814989

Type: Antibody

Proper Citation

Abcam Cat# ab188574, RRID:AB_2814989

Antibody Information

URL: http://antibodyregistry.org/AB_2814989

Proper Citation: Abcam Cat# ab188574, RRID:AB_2814989

Target Antigen: RAB8A

Host Organism: rabbit

Clonality: unknown

Comments: Applications: WB, IHC-P, ICC/IF, Flow Cyt

Antibody Name: Recombinant Anti-RAB8A antibody [EPR14873]

Description: This unknown targets RAB8A

Target Organism: rat, mouse, human

Clone ID: [EPR14873]

Antibody ID: AB_2814989

Vendor: Abcam

Catalog Number: ab188574

Record Creation Time: 20231110T032615+0000

Record Last Update: 20260829T055253+0000

Ratings and Alerts

No rating or validation information has been found for Recombinant Anti-RAB8A antibody [EPR14873].

No alerts have been found for Recombinant Anti-RAB8A antibody [EPR14873].

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 [RRID](#) .

Singh K, et al. (2026) Membrane Dysfunction as a Central Mechanism in LRRK2-Associated Parkinson's Disease: Comparative Analysis of G2019S and I1371V Variants. Cells, 15(4).

Shi W, et al. (2024) Targeting SHCBP1 Inhibits Tumor Progression by Restoring Ciliogenesis in Ductal Carcinoma. Cancer research, 84(24), 4156.

Abe T, et al. (2024) Lysosomal stress drives the release of pathogenic α -synuclein from macrophage lineage cells via the LRRK2-Rab10 pathway. iScience, 27(2), 108893.

Wang X, et al. (2023) Rab12 is a regulator of LRRK2 and its activation by damaged lysosomes. eLife, 12.

Bayati A, et al. (2022) Rapid macropinocytic transfer of α -synuclein to lysosomes. Cell reports, 40(3), 111102.

Scheiblich H, et al. (2021) Microglia jointly degrade fibrillar alpha-synuclein cargo by distribution through tunneling nanotubes. Cell, 184(20), 5089.

Kallemeijn WW, et al. (2021) Proteome-wide analysis of protein lipidation using chemical probes: in-gel fluorescence visualization, identification and quantification of N-myristoylation, N- and S-acylation, O-cholesterylation, S-farnesylation and S-geranylgeranylation. Nature protocols, 16(11), 5083.

Chen C, et al. (2020) Pathway-specific dysregulation of striatal excitatory synapses by LRRK2 mutations. eLife, 9.

Chen B, et al. (2018) Protein Lipidation in Cell Signaling and Diseases: Function, Regulation, and Therapeutic Opportunities. Cell chemical biology, 25(7), 817.