Resource Summary Report

Generated by NIF on Aug 29, 2026

Anti-GFP

RRID:AB_390913
Type: Antibody

Proper Citation

Roche Cat# 11814460001, RRID:AB_390913

Antibody Information

URL: http://antibodyregistry.org/AB_390913

Proper Citation: Roche Cat# 11814460001, RRID:AB_390913

Target Antigen: GFP

Host Organism: mouse

Clonality: monoclonal

Comments: Applications: Immunoprecipitation, Western blots, Immunostaining.

Antibody Name: Anti-GFP

Description: This monoclonal targets GFP

Clone ID: Clones 7.1 and 13.1

Defining Citation: PMID:20878781, PMID:19260068

Antibody ID: AB_390913

Vendor: Roche

Catalog Number: 11814460001

Record Creation Time: 20250820T030619+0000

Record Last Update: 20250820T152143+0000

Ratings and Alerts

No rating or validation information has been found for Anti-GFP.

No alerts have been found for Anti-GFP.

Data and Source Information

Source: Antibody Registry

Usage and Citation Metrics

We found 559 mentions in open access literature.

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

Bouton CR, et al. (2026) The long isoform of ZAP coordinates multiple enzymes to mediate complete decay of target transcripts. Cell reports, 45(5), 117305.

Sato K, et al. (2026) Alternative 3' UTRs and RNA-binding proteins Ewsr1b, HuR, and Syncrip organize sequential waves of translation to drive embryonic development. Cell reports, 45(6), 117557.

Abdrakhmanov A, et al. (2026) A lineage-specific selective autophagy receptor module mediates P-body turnover. Developmental cell, 61(4), 744.

Taskin AA, et al. (2026) Uncovering the initial response: Intra-mitochondrial surveillance activates the UPRmt. Molecular cell, 86(11), 2157.

Lemus L, et al. (2026) Lectins of the Emp47/ERGIC-53 family target misfolded O-mannosylated ER proteins to ER-associated degradation. Cell reports, 45(7), 117626.

Fischer JS, et al. (2026) A conserved mechanism of membrane fusion in nuclear pore complex assembly. Molecular cell.

Taneja N, et al. (2026) The actin crosslinker Fimbrin is required for force sensing at tricellular junctions. Developmental cell.

Xu L, et al. (2026) Longitudinal zone- and cell type-specific pattern-triggered immune responses in Arabidopsis roots. Developmental cell.

Qu W, et al. (2026) HRD1 negatively regulates autolysosome formation by inhibiting liquid-liquid phase separation of SNAP29. Cell reports, 45(2), 116914.

Kim SS, et al. (2026) Ribonuclease ? promotes longevity by preventing age-associated accumulation of circular RNA in stress granules. Molecular cell, 86(5), 885.

Zaremba E, et al. (2026) Vsb1, Ypq1, and Ypq2 control dynamic cationic amino acid storage in the yeast vacuole. Life science alliance, 9(7).

Chen Y, et al. (2026) Ribo-Tweezer: Rapid removal of ribosomal proteins reveals additional layers of post-transcriptional gene regulation. Molecular cell, 86(13), 2635.

Vind AC, et al. (2026) Mechanism of RACK1-dependent ZAK? activation at stalled and collided ribosomes. Molecular cell, 86(12), 2341.

Shao N, et al. (2026) Chloroplast sunscreening by protein condensates confers high-light tolerance. Cell.

Jiang M, et al. (2026) Butyrate causes loss of specific acetylation by the histone acetyltransferase p300. iScience, 29(7), 116528.

Lieb WS, et al. (2026) Protein kinase D deficiency induces a senescence-like phenotype in ?-cells and improves glucose and insulin tolerance under high-fat diet conditions. Molecular metabolism, 103, 102297.

Thomas V, et al. (2026) High-resolution mapping of the actin fusion focus reveals myosin V-dependent formin transport for aster formation. The Journal of cell biology, 225(2).

Álvarez D, et al. (2026) The molecular mechanism of lipid uptake by membrane-anchored bridge-like lipid transfer proteins. Proceedings of the National Academy of Sciences of the United States of America, 123(4), e2520399123.

Ye Y, et al. (2026) The MO25 protein Pmo25 functions in contractile ring stability and Sid2 localization during cytokinesis. iScience, 29(1), 114287.

Duhay V, et al. (2026) Control of lysosome function by the GTPase-activating protein TBC1D9B and its binding partner TMEM55B. Nature communications, 17(1).