Resource Summary Report

Generated by FDI Lab - SciCrunch.org on Apr 27, 2025

Mouse Anti-HA Monoclonal Antibody, Unconjugated, Clone 12CA5

RRID:AB_514505
Type: Antibody

Proper Citation

(Sigma-Aldrich Cat# 11583816001, RRID:AB_514505)

Antibody Information

URL: http://antibodyregistry.org/AB_514505

Proper Citation: (Sigma-Aldrich Cat# 11583816001, RRID:AB_514505)

Target Antigen: HA

Host Organism: mouse

Clonality: monoclonal

Comments: This antibody was sold by Roche, now it is sold by Sigma.

Antibody Name: Mouse Anti-HA Monoclonal Antibody, Unconjugated, Clone 12CA5

Description: This monoclonal targets HA

Clone ID: Clone 12CA5

Defining Citation: PMID:18425811

Antibody ID: AB_514505

Vendor: Sigma-Aldrich

Catalog Number: 11583816001

Record Creation Time: 20241016T234051+0000

Record Last Update: 20241017T010504+0000

Ratings and Alerts

No rating or validation information has been found for Mouse Anti-HA Monoclonal Antibody, Unconjugated, Clone 12CA5.

No alerts have been found for Mouse Anti-HA Monoclonal Antibody, Unconjugated, Clone 12CA5.

Data and Source Information

Source: Antibody Registry

Usage and Citation Metrics

We found 121 mentions in open access literature.

Listed below are recent publications. The full list is available at FDI Lab - SciCrunch.org.

Papadopoulos D, et al. (2024) The MYCN oncoprotein is an RNA-binding accessory factor of the nuclear exosome targeting complex. Molecular cell, 84(11), 2070.

Tofaute MJ, et al. (2024) SARS-CoV-2 NSP14 MTase activity is critical for inducing canonical NF-?B activation. Bioscience reports, 44(1).

Marmorale LJ, et al. (2024) Fast-evolving cofactors regulate the role of HEATR5 complexes in intra-Golgi trafficking. The Journal of cell biology, 223(3).

Hoh KL, et al. (2024) VAP-mediated membrane-tethering mechanisms implicate ER-PM contact function in pH homeostasis. Cell reports, 43(8), 114592.

Nakamori T, et al. (2024) The role of osteocrin in memory formation during early learning, as revealed by visual imprinting in chicks. iScience, 27(11), 111195.

Jeppsson K, et al. (2024) Loop-extruding Smc5/6 organizes transcription-induced positive DNA supercoils. Molecular cell, 84(5), 867.

Parnell EJ, et al. (2024) A conserved site on Ndc80 complex facilitates dynamic recruitment of Mps1 to yeast kinetochores to promote accurate chromosome segregation. Current biology : CB, 34(11), 2294.

Kunchala P, et al. (2024) Plasticity of the mitotic spindle in response to karyotype variation. Current biology : CB, 34(15), 3416.

Ruiz-Romero G, et al. (2024) Limiting 20S proteasome assembly leads to unbalanced nucleo-cytoplasmic distribution of 26S/30S proteasomes and chronic proteotoxicity. iScience, 27(11), 111095.

Kim DK, et al. (2023) A proteome-scale map of the SARS-CoV-2-human contactome. Nature biotechnology, 41(1), 140.

Valenti M, et al. (2023) Human gasdermin D and MLKL disrupt mitochondria, endocytic traffic and TORC1 signalling in budding yeast. Open biology, 13(5), 220366.

Marmorale LJ, et al. (2023) Two functionally distinct HEATR5 protein complexes are defined by fast-evolving co-factors in yeast. bioRxiv : the preprint server for biology.

Aguilar F, et al. (2023) Peptides from human BNIP5 and PXT1 and non-native binders of pro-apoptotic BAK can directly activate or inhibit BAK-mediated membrane permeabilization. Structure (London, England : 1993), 31(3), 265.

Oppenheim T, et al. (2023) The Cdc48 N-terminal domain has a molecular switch that mediates the Npl4-Ufd1-Cdc48 complex formation. Structure (London, England : 1993), 31(7), 764.

Wu X, et al. (2023) Secreted ORF8 induces monocytic pro-inflammatory cytokines through NLRP3 pathways in patients with severe COVID-19. iScience, 26(6), 106929.

Guichard A, et al. (2023) A comprehensive Drosophila resource to identify key functional interactions between SARS-CoV-2 factors and host proteins. Cell reports, 42(8), 112842.

Prieto-Ruiz F, et al. (2023) Myosin II regulatory light chain phosphorylation and formin availability modulate cytokinesis upon changes in carbohydrate metabolism. eLife, 12.

Neumann H, et al. (2023) Ratcheted transport and sequential assembly of the yeast telomerase RNP. Cell reports, 42(12), 113565.

Hsieh WC, et al. (2022) Glucose starvation induces a switch in the histone acetylome for activation of gluconeogenic and fat metabolism genes. Molecular cell, 82(1), 60.

Sun W, et al. (2022) GFI1 Cooperates with IKZF1/IKAROS to Activate Gene Expression in T-cell Acute Lymphoblastic Leukemia. Molecular cancer research : MCR, 20(4), 501.