

Resource Summary Report

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Mouse Anti-HA tag Monoclonal Antibody, Unconjugated, Clone HA.C5

RRID:AB_444303

Type: Antibody

Proper Citation

(Abcam Cat# ab18181, RRID:AB_444303)

Antibody Information

URL: http://antibodyregistry.org/AB_444303

Proper Citation: (Abcam Cat# ab18181, RRID:AB_444303)

Target Antigen: HA tag

Host Organism: mouse

Clonality: monoclonal

Comments: validation status unknown, seller recommendations provided in 2012: ELISA; Immunocytochemistry; Immunofluorescence; Immunoprecipitation; Western Blot; Immunocytochemistry, Immunocytochemistry/Immunofluorescence, Immunoprecipitation, Western Blot

Antibody Name: Mouse Anti-HA tag Monoclonal Antibody, Unconjugated, Clone HA.C5

Description: This monoclonal targets HA tag

Clone ID: Clone HA.C5

Antibody ID: AB_444303

Vendor: Abcam

Catalog Number: ab18181

Record Creation Time: 20241016T232707+0000

Record Last Update: 20241017T004248+0000

Ratings and Alerts

No rating or validation information has been found for Mouse Anti-HA tag Monoclonal Antibody, Unconjugated, Clone HA.C5.

No alerts have been found for Mouse Anti-HA tag Monoclonal Antibody, Unconjugated, Clone HA.C5.

Data and Source Information

Source: [Antibody Registry](#)

Usage and Citation Metrics

We found 43 mentions in open access literature.

Listed below are recent publications. The full list is available at [FDI Lab - SciCrunch.org](#).

Hou SS, et al. (2024) Recording γ -secretase activity in living mouse brains. eLife, 13.

Smirnova AM, et al. (2024) Stem-loop-induced ribosome queuing in the uORF2/ATF4 overlap fine-tunes stress-induced human ATF4 translational control. Cell reports, 43(4), 113976.

Qin Y, et al. (2024) TRIM37 is a primate-specific E3 ligase for Huntingtin and accounts for the striatal degeneration in Huntington's disease. Science advances, 10(20), ead12036.

Smirnova AM, et al. (2024) Protocol for detecting and quantifying hATF4-HA in non-stress versus stress conditions using automated and quantitative Jess western blotting. STAR protocols, 5(4), 103454.

Layden HM, et al. (2024) Mutant FOXO1 controls an oncogenic network via enhancer accessibility. Cell genomics, 4(4), 100537.

Pal D, et al. (2023) Mutating novel interaction sites in NRP1 reduces SARS-CoV-2 spike protein internalization. iScience, 26(4), 106274.

Kschonsak YT, et al. (2023) Potent and selective binders of the E3 ubiquitin ligase ZNRF3 stimulate Wnt signaling and intestinal organoid growth. Cell chemical biology.

van Solinge TS, et al. (2023) Illuminating cellular and extracellular vesicle-mediated communication via a split-Nanoluc reporter in vitro and in vivo. Cell reports methods, 3(2), 100412.

Jiang R, et al. (2023) TLR7 neo-functionalizes to sense dsRNA and trigger antiviral and antibacterial immunity in non-tetrapod vertebrates. *iScience*, 26(12), 108315.

Zhu Z, et al. (2023) Development of a DNA aptamer targeting IDO1 with anti-tumor effects. *iScience*, 26(8), 107367.

Thomasen PB, et al. (2023) SorCS2 binds progranulin to regulate motor neuron development. *Cell reports*, 42(11), 113333.

Wang N, et al. (2022) Inactivation of a wheat protein kinase gene confers broad-spectrum resistance to rust fungi. *Cell*, 185(16), 2961.

Coillard L, et al. (2022) The NFAT3/RERG Complex in Luminal Breast Cancers Is Required to Inhibit Cell Invasion and May Be Correlated With an Absence of Axillary Lymph Nodes Colonization. *Frontiers in oncology*, 12, 804868.

Ding P, et al. (2022) Intracellular complement C5a/C5aR1 stabilizes β -catenin to promote colorectal tumorigenesis. *Cell reports*, 39(9), 110851.

Melani R, et al. (2022) Inhibitory co-transmission from midbrain dopamine neurons relies on presynaptic GABA uptake. *Cell reports*, 39(3), 110716.

Qi H, et al. (2022) The ADP-ribose hydrolase NUDT5 is important for DNA repair. *Cell reports*, 41(12), 111866.

Smutná T, et al. (2022) A cytosolic ferredoxin-independent hydrogenase possibly mediates hydrogen uptake in *Trichomonas vaginalis*. *Current biology : CB*, 32(1), 124.

Fiskus W, et al. (2022) Effective Menin inhibitor-based combinations against AML with MLL rearrangement or NPM1 mutation (NPM1c). *Blood cancer journal*, 12(1), 5.

Xiao H, et al. (2022) Architecture of the outbred brown fat proteome defines regulators of metabolic physiology. *Cell*, 185(24), 4654.

Tomasello DL, et al. (2022) 16pdel lipid changes in iPSC-derived neurons and function of FAM57B in lipid metabolism and synaptogenesis. *iScience*, 25(1), 103551.