

Resource Summary Report

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Influenza hemagglutinin epitope (YPYDVPDYA)

RRID:AB_2314672

Type: Antibody

Proper Citation

Covance Cat# MMS-101P, RRID:AB_2314672

Antibody Information

URL: http://antibodyregistry.org/AB_2314672

Proper Citation: Covance Cat# MMS-101P, RRID:AB_2314672

Clonality: unknown

Antibody Name: Influenza hemagglutinin epitope (YPYDVPDYA)

Description: This unknown targets

Defining Citation: [PMID:19235225](#)

Antibody ID: AB_2314672

Vendor: Covance

Catalog Number: MMS-101P

Record Creation Time: 20231110T042044+0000

Record Last Update: 20260829T043521+0000

Ratings and Alerts

- ENCODE PROJECT External validation for lot: unknown is available under ENCODE ID: ENCAB000ASV - ENCODE <https://www.encodeproject.org/antibodies/ENCAB000ASV>

No alerts have been found for Influenza hemagglutinin epitope (YPYDVPDYA).

Data and Source Information

Usage and Citation Metrics

We found 107 mentions in open access literature.

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

Vo N, et al. (2026) Detection of nuclear STING in cultured human cells and in the normal and cancer tissues. *iScience*, 29(5), 115711.

Luo Y, et al. (2026) mRNA 3' UTRs chaperone intrinsically disordered regions to control protein activity. *Cell*, 189(15), 4619.

Latini C, et al. (2025) Cytoplasmic DIS3 is an exosome-independent endoribonuclease with catalytic activity toward circular RNAs. *Cell reports*, 44(6), 115769.

Al Kabbani MA, et al. (2025) Effects of P301L-TAU on post-translational modifications of microtubules in human iPSC-derived cortical neurons and TAU transgenic mice. *Neural regeneration research*, 20(8), 2348.

Sahasrabuddhe AA, et al. (2025) The FBXO45-GEF-H1 Axis Controls Germinal Center Formation and B-cell Lymphomagenesis. *Cancer discovery*, 15(4), 838.

Jeong HK, et al. (2025) Micro-scale control of oligodendrocyte morphology and myelination by the intellectual disability-linked protein acyltransferase ZDHHC9. *eLife*, 13.

Xu B, et al. (2025) γ -synuclein blocks γ -synuclein condensate fusion to disrupt the maturation of phase separation. *Cell reports*, 44(6), 115761.

Wang J, et al. (2024) ARF alters PAF1 complex integrity to selectively repress oncogenic transcription programs upon p53 loss. *Molecular cell*, 84(23), 4538.

Anderson MJM, et al. (2024) Molecular basis of proteolytic cleavage regulation by the extracellular matrix receptor dystroglycan. *Structure (London, England : 1993)*.

Chen L, et al. (2024) Palmitoylation alters LDHA activity and pancreatic cancer response to chemotherapy. *Cancer letters*, 587, 216696.

Rona G, et al. (2024) CDK-independent role of D-type cyclins in regulating DNA mismatch repair. *Molecular cell*.

Le Guerroué F, et al. (2023) TNIP1 inhibits selective autophagy via bipartite interaction with LC3/GABARAP and TAX1BP1. *Molecular cell*, 83(6), 927.

Dardis GJ, et al. (2023) An EZH2-NF- κ B regulatory axis drives expression of pro-oncogenic gene signatures in triple negative breast cancer. *iScience*, 26(7), 107115.

Song T, et al. (2023) TRIM28 represses renal cell carcinoma cell proliferation by inhibiting TFE3/KDM6A-regulated autophagy. *The Journal of biological chemistry*, 299(5), 104621.

Mishra I, et al. (2022) Protein tyrosine phosphatase receptor ? serves as the orexigenic asprosin receptor. *Cell metabolism*, 34(4), 549.

Koester AM, et al. (2022) EFR3 and phosphatidylinositol 4-kinase III? regulate insulin-stimulated glucose transport and GLUT4 dispersal in 3T3-L1 adipocytes. *Bioscience reports*, 42(7).

Lv S, et al. (2022) Efficient and reversible Cas13d-mediated knockdown with an all-in-one lentivirus-vector. *Frontiers in bioengineering and biotechnology*, 10, 960192.

Shi DD, et al. (2022) De novo pyrimidine synthesis is a targetable vulnerability in IDH mutant glioma. *Cancer cell*, 40(9), 939.

Huppertz I, et al. (2022) Riboregulation of Enolase 1 activity controls glycolysis and embryonic stem cell differentiation. *Molecular cell*, 82(14), 2666.

Marshall RS, et al. (2022) A trio of ubiquitin ligases sequentially drives ubiquitylation and autophagic degradation of dysfunctional yeast proteasomes. *Cell reports*, 38(11), 110535.