

Resource Summary Report

Generated by [NIF](#) on Jul 30, 2026

Anti-His Tag, clone HIS.H8

RRID:AB_492660

Type: Antibody

Proper Citation

Millipore Cat# 05-949, RRID:AB_492660

Antibody Information

URL: http://antibodyregistry.org/AB_492660

Proper Citation: Millipore Cat# 05-949, RRID:AB_492660

Target Antigen: His Tag clone HIS.H8

Host Organism: mouse

Clonality: monoclonal

Comments: seller recommendations: IgG2; IgG2 IC, IP, WB; Western Blot; Immunocytochemistry; Immunoprecipitation

Antibody Name: Anti-His Tag, clone HIS.H8

Description: This monoclonal targets His Tag clone HIS.H8

Target Organism: all

Antibody ID: AB_492660

Vendor: Millipore

Catalog Number: 05-949

Record Creation Time: 20250819T213605+0000

Record Last Update: 20250819T232933+0000

Ratings and Alerts

No rating or validation information has been found for Anti-His Tag, clone HIS.H8.

No alerts have been found for Anti-His Tag, clone HIS.H8.

Data and Source Information

Source: [Antibody Registry](#)

Usage and Citation Metrics

We found 12 mentions in open access literature.

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

Trofimova D, et al. (2025) Fungal kinesin-8 motors dimerize by folding their proximal tail domain into a compact helical bundle. *Structure* (London, England : 1993).

Moura M, et al. (2025) Polo kinase inhibits protein phosphatase 1 to promote the spindle assembly checkpoint and prevent aneuploidy. *Current biology : CB*, 35(21), 5289.

Heisler DB, et al. (2023) A concerted mechanism involving ACAT and SREBPs by which oxysterols deplete accessible cholesterol to restrict microbial infection. *eLife*, 12.

Lebedev M, et al. (2023) Anillin forms linear structures and facilitates furrow ingression after septin and formin depletion. *Cell reports*, 42(9), 113076.

Datta S, et al. (2023) High-resolution photocatalytic mapping of SARS-CoV-2 spike interactions on the cell surface. *Cell chemical biology*, 30(10), 1313.

Seraphim TV, et al. (2022) Assembly principles of the human R2TP chaperone complex reveal the presence of R2T and R2P complexes. *Structure* (London, England : 1993), 30(1), 156.

Cuevas-Navarro A, et al. (2021) The RAS GTPase RIT1 compromises mitotic fidelity through spindle assembly checkpoint suppression. *Current biology : CB*, 31(17), 3915.

Yoon S, et al. (2020) TGF- β -Induced Phosphorylation of Usp9X Stabilizes Ankyrin-G and Regulates Dendritic Spine Development and Maintenance. *Cell reports*, 31(8), 107685.

Ko HL, et al. (2019) HNF4 β Combinatorial Isoform Heterodimers Activate Distinct Gene Targets that Differ from Their Corresponding Homodimers. *Cell reports*, 26(10), 2549.

Endapally S, et al. (2019) Molecular Discrimination between Two Conformations of Sphingomyelin in Plasma Membranes. *Cell*, 176(5), 1040.

Beuret N, et al. (2017) Amyloid-like aggregation of provasopressin in diabetes insipidus and secretory granule sorting. *BMC biology*, 15(1), 5.

Arellano-Santoyo H, et al. (2017) A Tubulin Binding Switch Underlies Kip3/Kinesin-8 Depolymerase Activity. *Developmental cell*, 42(1), 37.