

Resource Summary Report

Generated by [FDI Lab - SciCrunch.org](#) on Apr 18, 2025

OLLAS Epitope Tag Antibody (L2) - BSA Free

RRID:AB_1625979

Type: Antibody

Proper Citation

(Novus Cat# NBP1-06713, RRID:AB_1625979)

Antibody Information

URL: http://antibodyregistry.org/AB_1625979

Proper Citation: (Novus Cat# NBP1-06713, RRID:AB_1625979)

Target Antigen: OLLAS Epitope Tag

Host Organism: Rat

Clonality: monoclonal

Comments: Applications: Western Blot, Immunohistochemistry, Immunocytochemistry/Immunofluorescence, Immunoprecipitation, Immunohistochemistry-Paraffin, Immunohistochemistry-Frozen, Immunoblotting, Immunohistochemistry Whole-Mount, Chromatin Immunoprecipitation Sequencing

Antibody Name: OLLAS Epitope Tag Antibody (L2) - BSA Free

Description: This monoclonal targets OLLAS Epitope Tag

Target Organism: Epitope Tag

Clone ID: L2

Antibody ID: AB_1625979

Vendor: Novus

Catalog Number: NBP1-06713

Alternative Catalog Numbers: NBP1-06713SS

Record Creation Time: 20241016T230828+0000

Record Last Update: 20241017T000642+0000

Ratings and Alerts

No rating or validation information has been found for OLLAS Epitope Tag Antibody (L2) - BSA Free.

No alerts have been found for OLLAS Epitope Tag Antibody (L2) - BSA Free.

Data and Source Information

Source: [Antibody Registry](#)

Usage and Citation Metrics

We found 10 mentions in open access literature.

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

Sanfilippo P, et al. (2024) Mapping of multiple neurotransmitter receptor subtypes and distinct protein complexes to the connectome. *Neuron*, 112(6), 942.

Gleason RJ, et al. (2023) Developmentally programmed histone H3 expression regulates cellular plasticity at the parental-to-early embryo transition. *Science advances*, 9(14), eadh0411.

Sanfilippo P, et al. (2023) Mapping of multiple neurotransmitter receptor subtypes and distinct protein complexes to the connectome. *bioRxiv : the preprint server for biology*.

Bauer M, et al. (2023) Heterodimerization-dependent secretion of bone morphogenetic proteins in *Drosophila*. *Developmental cell*, 58(8), 645.

Jeon H, et al. (2022) Topographic connectivity and cellular profiling reveal detailed input pathways and functionally distinct cell types in the subthalamic nucleus. *Cell reports*, 38(9), 110439.

Schmidt H, et al. (2021) Protein-based condensation mechanisms drive the assembly of RNA-rich P granules. *eLife*, 10.

Robinson-Thiewes S, et al. (2021) Non-autonomous regulation of germline stem cell proliferation by somatic MPK-1/MAPK activity in *C. elegans*. *Cell reports*, 35(8), 109162.

Lee CS, et al. (2020) Recruitment of mRNAs to P granules by condensation with intrinsically-disordered proteins. *eLife*, 9.

Akay A, et al. (2017) The Helicase Aquarius/EMB-4 Is Required to Overcome Intronic Barriers to Allow Nuclear RNAi Pathways to Heritably Silence Transcription. *Developmental cell*, 42(3), 241.

Lee CS, et al. (2017) Nanos promotes epigenetic reprogramming of the germline by down-regulation of the THAP transcription factor LIN-15B. *eLife*, 6.