

Resource Summary Report

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Anti-digoxigenin

RRID:AB_514496

Type: Antibody

Proper Citation

(Sigma-Aldrich Cat# 11333089001, RRID:AB_514496)

Antibody Information

URL: http://antibodyregistry.org/AB_514496

Proper Citation: (Sigma-Aldrich Cat# 11333089001, RRID:AB_514496)

Target Antigen: digoxinenin

Host Organism: sheep

Clonality: polyclonal

Comments: This is a Roche product, offered by Sigma.

Antibody Name: Anti-digoxigenin

Description: This polyclonal targets digoxinenin

Antibody ID: AB_514496

Vendor: Sigma-Aldrich

Catalog Number: 11333089001

Record Creation Time: 20231110T034746+0000

Record Last Update: 20240725T011342+0000

Ratings and Alerts

No rating or validation information has been found for Anti-digoxigenin.

No alerts have been found for Anti-digoxigenin.

Data and Source Information

Source: [Antibody Registry](#)

Usage and Citation Metrics

We found 29 mentions in open access literature.

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

Ku J, et al. (2024) Alternative polyadenylation determines the functional landscape of inverted Alu repeats. *Molecular cell*.

Kasal MR, et al. (2023) Lon degrades stable substrates slowly but with enhanced processivity, redefining the attributes of a successful AAA+ protease. *Cell reports*, 42(9), 113061.

Horio T, et al. (2023) Regulation of RNG105/caprin1 dynamics by pathogenic cytoplasmic FUS and TDP-43 in neuronal RNA granules modulates synaptic loss. *Heliyon*, 9(6), e17065.

Galupa R, et al. (2023) Enhancer architecture and chromatin accessibility constrain phenotypic space during *Drosophila* development. *Developmental cell*, 58(1), 51.

McKie SJ, et al. (2022) Topoisomerase VI is a chirally-selective, preferential DNA decatenase. *eLife*, 11.

Cui H, et al. (2022) Protocol to image and quantify nucleocytoplasmic transport in cultured cells using fluorescent in situ hybridization and a dual reporter system. *STAR protocols*, 3(4), 101813.

Gandara L, et al. (2022) Developmental phenomics suggests that H3K4 monomethylation confers multi-level phenotypic robustness. *Cell reports*, 41(11), 111832.

Janissen R, et al. (2022) High-throughput single-molecule experiments reveal heterogeneity, state switching, and three interconnected pause states in transcription. *Cell reports*, 39(4), 110749.

Kuijpers L, et al. (2022) Characterizing single-molecule dynamics of viral RNA-dependent RNA polymerases with multiplexed magnetic tweezers. *STAR protocols*, 3(3), 101606.

Janissen R, et al. (2021) Induced intra- and intermolecular template switching as a therapeutic mechanism against RNA viruses. *Molecular cell*, 81(21), 4467.

Domsch K, et al. (2021) The Hox Transcription Factor Ubx Ensures Somatic Myogenesis by Suppressing the Mesodermal Master Regulator Twist. *Cell reports*, 34(1), 108577.

Bera SC, et al. (2021) The nucleotide addition cycle of the SARS-CoV-2 polymerase. *Cell reports*, 36(9), 109650.

Kurihara M, et al. (2020) Genomic Profiling by ALaP-Seq Reveals Transcriptional Regulation by PML Bodies through DNMT3A Exclusion. *Molecular cell*, 78(3), 493.

Fukuda S, et al. (2020) The Biogenesis of SRP RNA Is Modulated by an RNA Folding Intermediate Attained during Transcription. *Molecular cell*, 77(2), 241.

Djabrayan NJ, et al. (2019) Metabolic Regulation of Developmental Cell Cycles and Zygotic Transcription. *Current biology : CB*, 29(7), 1193.

Johnson HE, et al. (2019) Signaling Dynamics Control Cell Fate in the Early *Drosophila* Embryo. *Developmental cell*, 48(3), 361.

Beier KT, et al. (2019) Topological Organization of Ventral Tegmental Area Connectivity Revealed by Viral-Genetic Dissection of Input-Output Relations. *Cell reports*, 26(1), 159.

Seol Y, et al. (2019) Homology sensing via non-linear amplification of sequence-dependent pausing by RecQ helicase. *eLife*, 8.

Ohno M, et al. (2019) Sub-nucleosomal Genome Structure Reveals Distinct Nucleosome Folding Motifs. *Cell*, 176(3), 520.

Le TT, et al. (2019) Synergistic Coordination of Chromatin Torsional Mechanics and Topoisomerase Activity. *Cell*, 179(3), 619.