

Resource Summary Report

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Diagenode

RRID:SCR_014807

Type: Tool

Proper Citation

Diagenode (RRID:SCR_014807)

Resource Information

URL: <https://www.diagenode.com>

Proper Citation: Diagenode (RRID:SCR_014807)

Description: Commercial provider of equipment and resources for epigenetics research, biological sample preparation, and diagnostic assays. The products Diagenode provides include automation and semi-automation software, antibodies, reagents, kits, and shearing technologies.

Resource Type: antibody supplier, material resource, reagent supplier

Keywords: epigenetics, sample preparation system, diagnostic device, reagent, assay, antibody supplier, instrument supplier, methylation, cell culture, automated, shearing

Resource Name: Diagenode

Resource ID: SCR_014807

Record Creation Time: 20220129T080322+0000

Record Last Update: 20260829T182855+0000

Ratings and Alerts

No rating or validation information has been found for Diagenode.

No alerts have been found for Diagenode.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 985 mentions in open access literature.

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

Cortez J, et al. (2026) Reference genome of an irruptive migrant, the pine siskin (*Spinus pinus*). *The Journal of heredity*, 117(3), 469.

Carlson BJ, et al. (2026) A reference genome assembly for the pink volcano barnacle, *Tetraclita rubescens*, Nilsson-Cantell, 1931. *The Journal of heredity*, 117(1), 173.

Hallal R, et al. (2026) Targeting SUMOylation Triggers IFN- γ -dependent Activation of Patient and Allogenic NK Cells in Preclinical Models of Acute Myeloid Leukemia. *Molecular cancer therapeutics*, 25(1), 125.

Woods RM, et al. (2026) Developmental modulation of schizophrenia risk gene methylation in offspring exhibiting cognitive deficits following maternal immune activation. *Molecular psychiatry*, 31(1), 418.

Hallas JM, et al. (2026) A genome assembly for mule deer, *Odocoileus hemionus*, from Southern California. *The Journal of heredity*, 117(3), 502.

Rivera MG, et al. (2026) Genome assembly and annotation of a deep-diving pinniped, the northern elephant seal (*Mirounga angustirostris*). *The Journal of heredity*, 117(3), 512.

Richmond JQ, et al. (2026) A chromosome-level reference genome assembly for Gilbert's skink *Plestiodon gilberti*. *The Journal of heredity*, 117(1), 115.

Zeng C, et al. (2026) A SET domain-containing protein and HCF-1 maintain transgenerational epigenetic memory. *Nature communications*, 17(1), 1462.

Smith MG, et al. (2026) Hypoxic stress granules trigger immunogenic dormancy in lung cancer. *bioRxiv : the preprint server for biology*.

Alsolami S, et al. (2026) Stable bioreactor control reveals acidic pH-driven metabolic reprogramming and mitochondrial dysfunction in human lymphoblastoid cells. *Communications biology*, 9(1).

Turpin M, et al. (2026) TDG orchestrates ATF4-dependent gene transcription during retinoic acid-induced cell fate acquisition. *Nucleic acids research*, 54(5).

Han Y, et al. (2026) Glycine max Jumonji C domain-containing proteins JM19/20 exhibit endopeptidase activity and interact with LUXs to mediate flowering time. *Nucleic acids research*, 54(5).

Oosterlynck M, et al. (2026) Tau profiling of brain extracellular vesicles reveals PHF6 peptide as core for pathological tau seeding in Alzheimer's disease. *Journal of biomedical science*, 33(1).

Bournaud C, et al. (2026) Profiling DNA-protein interactions in *Meloidogyne incognita* using dCas9-based affinity purification. *Plant methods*, 22(1), 20.

Wutthituntisil N, et al. (2026) A reference genome for the rough limpet *Lottia Scabra*, an intertidal species from the northeastern Pacific. *The Journal of heredity*, 117(2), 299.

Centonze A, et al. (2026) A Plastic EMP1+ to LGR5+ Cell State Conversion as a Bypass to KRASG12D Pharmacologic Inhibition in Metastatic Colorectal Cancer. *Cancer discovery*, 16(2), 320.

Kama Y, et al. (2026) Notch interaction with RUNX factors regulates initiation of the T-lineage program. *The Journal of experimental medicine*, 223(2).

Hjelmgren L, et al. (2026) Dysregulation of the DNA damage response by phosphorothioate antisense oligonucleotides. *Nature communications*, 17(1).

Hess A, et al. (2026) Non-disruptive in vitro monitoring of cellular states with cell-free DNA methylation. *Genome biology*, 27(1).

Desgraupes S, et al. (2026) The genetic driver of Acute Necrotizing Encephalopathy, RANBP2, regulates the inflammatory response to Influenza A virus infection. *Nature communications*, 17(1).