

Resource Summary Report

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LabArchives

RRID:SCR_013973

Type: Tool

Proper Citation

LabArchives (RRID:SCR_013973)

Resource Information

URL: <http://www.labarchives.com>

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Description: An ELN (electronic lab notebook) software application where researchers can record and organize their data in lieu of a paper notebook. Users can store and edit texts, PDFs, spreadsheets, images, sample collections, and other types of data. LabArchives automatically backs up data and data can be accessed anywhere. Users can choose to use the professional edition of LabArchives, the classroom edition, or to purchase an enterprise license.

Resource Type: software application, software resource, electronic laboratory notebook

Keywords: ELN, electronic laboratory notebook, software application, data management, lab management, professional, classroom, enterprise

Funding:

Availability: Available to the research and education community

Resource Name: LabArchives

Resource ID: SCR_013973

Record Creation Time: 20220129T080318+0000

Record Last Update: 20250410T070504+0000

Ratings and Alerts

No rating or validation information has been found for LabArchives.

No alerts have been found for LabArchives.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 60 mentions in open access literature.

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

Jhaji KS, et al. (2024) Use of Artificial Intelligence Tools for Research by Medical Students: A Narrative Review. *Cureus*, 16(3), e55367.

Bourke J, et al. (2024) Efficacy of heel lifts for mid-portion Achilles tendinopathy (the LIFT trial): study protocol for a randomised controlled trial. *Trials*, 25(1), 345.

Francis EK, et al. (2023) Refugia or reservoir? Feral goats and their role in the maintenance and circulation of benzimidazole-resistant gastrointestinal nematodes on shared pastures. *Parasitology*, 150(8), 672.

Baláž V, et al. (2023) Angie-LAMP for diagnosis of human eosinophilic meningitis using dog as proxy: A LAMP assay for *Angiostrongylus cantonensis* DNA in cerebrospinal fluid. *PLoS neglected tropical diseases*, 17(5), e0011038.

Berezin CT, et al. (2023) Ten simple rules for managing laboratory information. *PLoS computational biology*, 19(12), e1011652.

Forst DA, et al. (2023) Study protocol for NeuroCARE: a randomised controlled trial of a psychological intervention for caregivers of patients with primary malignant brain tumours. *BMJ open*, 13(9), e069410.

Power RI, et al. (2022) Exploration of the sensitivity to macrocyclic lactones in the canine heartworm (*Dirofilaria immitis*) in Australia using phenotypic and genotypic approaches. *International journal for parasitology. Drugs and drug resistance*, 20, 145.

Harris V, et al. (2022) Healthy Mothers Healthy Families, Health Promoting Activity Coaching for mothers of children with a disability: Exploring mothers' perspectives of programme feasibility. *Australian occupational therapy journal*, 69(6), 662.

Rivory P, et al. (2020) Apparent lack of spill-over of parasites from an invasive anuran: PCR detects *Entamoeba* in cane toads (*Rhinella marina*) but not in sympatric Australian native frogs. *International journal for parasitology. Parasites and wildlife*, 12, 207.

Chandra S, et al. (2020) Biotic Factors Influence Microbiota of Nymph Ticks from Vegetation in Sydney, Australia. *Pathogens (Basel, Switzerland)*, 9(7).

Andersson MI, et al. (2020) SARS-CoV-2 RNA detected in blood products from patients with COVID-19 is not associated with infectious virus. *Wellcome open research*, 5, 181.

Tenopir C, et al. (2020) Data sharing, management, use, and reuse: Practices and perceptions of scientists worldwide. *PLoS one*, 15(3), e0229003.

González R, et al. (2019) Mutagenesis Scanning Uncovers Evolutionary Constraints on Tobacco Etch Potyvirus Membrane-Associated 6K2 Protein. *Genome biology and evolution*, 11(4), 1207.

Sacco A, et al. (2019) New insights in the control of antioxidants accumulation in tomato by transcriptomic analyses of genotypes exhibiting contrasting levels of fruit metabolites. *BMC genomics*, 20(1), 43.

Ambrós S, et al. (2018) Engineered Functional Redundancy Relaxes Selective Constraints upon Endogenous Genes in Viral RNA Genomes. *Genome biology and evolution*, 10(7), 1823.

Carbonell P, et al. (2018) An automated Design-Build-Test-Learn pipeline for enhanced microbial production of fine chemicals. *Communications biology*, 1, 66.

Koufariotis L, et al. (2018) Sequencing the mosaic genome of Brahman cattle identifies historic and recent introgression including polled. *Scientific reports*, 8(1), 17761.

Neuhaus K, et al. (2017) Differentiation of ncRNAs from small mRNAs in *Escherichia coli* O157:H7 EDL933 (EHEC) by combined RNAseq and RIBOseq - *ryhB* encodes the regulatory RNA *RyhB* and a peptide, *RyhP*. *BMC genomics*, 18(1), 216.

Shikha M, et al. (2017) Genomic Selection for Drought Tolerance Using Genome-Wide SNPs in Maize. *Frontiers in plant science*, 8, 550.

Riley EM, et al. (2017) Implementation and use of cloud-based electronic lab notebook in a bioprocess engineering teaching laboratory. *Journal of biological engineering*, 11, 40.