

Resource Summary Report

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Protocols.io

RRID:SCR_010490

Type: Tool

Proper Citation

Protocols.io (RRID:SCR_010490)

Resource Information

URL: <http://protocols.io/>

Proper Citation: Protocols.io (RRID:SCR_010490)

Description: A commercial protocol sharing tool, which allows scientists to check in protocols the way that computer scientists check in code.

Resource Type: commercial organization, software resource

Resource Name: Protocols.io

Resource ID: SCR_010490

Alternate IDs: nlx_158033

Record Creation Time: 20220129T080259+0000

Record Last Update: 20260905T133344+0000

Ratings and Alerts

No rating or validation information has been found for Protocols.io.

No alerts have been found for Protocols.io.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 267 mentions in open access literature.

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

Hill JL, et al. (2026) Transcriptomic profiling of the embryonic *C. elegans* intestine with single-cell resolution. *bioRxiv : the preprint server for biology*.

Byrne J, et al. (2026) Single-nucleus profiling reveals age-associated remodeling opposed by parity in the postmenopausal human ovary. *bioRxiv : the preprint server for biology*.

Larue L, et al. (2026) A Short Report on Melanocyte/Melanoma Culture, Senescence, and Reproducibility. *Pigment cell & melanoma research*, 39(2), e70084.

Dautan D, et al. (2026) Gut-initiated alpha synuclein fibrils drive parkinsonism phenotypes: temporal mapping of REM sleep behavior disorder-like and other non-motor symptoms. *Translational neurodegeneration*, 15(1).

Liu J, et al. (2026) Mapping the specificity of H3N2 strain-specific and cross-reactive human neutralizing antibodies elicited by the 2025-2026 influenza vaccine. *medRxiv : the preprint server for health sciences*.

Naing SY, et al. (2026) Impact of reducing colistin use on colistin resistance in *Escherichia coli* isolated from humans and poultry in Indonesia (COINCIDE): A protocol for a multisectoral, transdisciplinary One Health study. *One health (Amsterdam, Netherlands)*, 22, 101347.

Chowdhury S, et al. (2026) Conserved HSP60 structure with lineage- and context-specific regulation in cnidarians. *Life science alliance*, 9(9).

Amador LG, et al. (2026) Bridging data silos to holistically model plant macrophenology. *The New phytologist*, 251(2), 625.

Surles-Zeigler MC, et al. (2026) PRECISE TBI Model Catalog: Increasing Accessibility and Reproducibility in TBI research. *bioRxiv : the preprint server for biology*.

Gillman AN, et al. (2026) EZ-DMS - A Simple and Accessible Protocol and Software Package for Deep Mutational Scanning of Virus Proteins. *bioRxiv : the preprint server for biology*.

Rohilla R, et al. (2026) Molecular Epidemiology of *Escherichia coli* Clinical Isolates From a Tertiary Care Hospital in Uttarakhand, India: Insights Into Antimicrobial Resistance (AMR) Burden, Virulence Profiles, and High-Risk Lineages. *Cureus*, 18(1), e102613.

Obermayer A, et al. (2026) A block staining method using ethanolic phosphotungstic acid for the visualisation of collagens in transmission electron microscopy. *PloS one*, 21(2), e0339342.

Klepper A, et al. (2026) Immune profiling links autoimmune hepatitis to human herpesvirus 6 and relaxin receptor antigens. *The Journal of experimental medicine*, 223(4).

Telles-de-Deus J, et al. (2026) Evolution and spillover dynamics of yellow fever at the forest-urban interface in Brazil. *Nature microbiology*, 11(4), 877.

Heinl C, et al. (2026) A call for preregistration of in vitro research : Preregistration as a tool to improve reproducibility and transparency. *EMBO reports*, 27(9), 2147.

Ulloa GM, et al. (2026) Zoonotic and Anthroponotic *Plasmodium* spp. Circulation between Wild Primates and Indigenous Community, Peruvian Amazon, 2007-2020. *Emerging infectious diseases*, 32(5), 707.

Fisher SA, et al. (2026) The HuBMAP Framework for Advancing Data FAIRness. *bioRxiv* : the preprint server for biology.

Bourgikos E, et al. (2026) Evolutionary history of Jamestown Canyon virus disentangles complex multi-vector ecology. *bioRxiv* : the preprint server for biology.

Pan WJ, et al. (2026) An integrated platform for simultaneous wide-field voltage/calcium imaging and fMRI (EPI & ZTE) reveals neuronal infraslow dynamics underlying functional connectivity. *bioRxiv* : the preprint server for biology.

Knebel E, et al. (2026) Construction of saturated Tn-Seq libraries of *Brucella abortus* S19 for transposon insertion and effective density analysis across stress conditions. *Microbiology resource announcements*, 15(5), e0109925.