

Resource Summary Report

Generated by [kravitz2](#) on Jul 31, 2026

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: software resource, commercial organization

Resource Name: Protocols.io

Resource ID: SCR_010490

Alternate IDs: nlx_158033

Record Creation Time: 20220129T080259+0000

Record Last Update: 20260731T042833+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 198 mentions in open access literature.

Listed below are recent publications. The full list is available at [kravitz2](#) .

Heuston EF, et al. (2025) Optimized methods for scRNA-seq and snRNA-seq of skeletal muscle stored in nucleic acid stabilizing preservative. *Communications biology*, 8(1), 10.

Wright A, et al. (2025) Glial activation and nociceptive neuropeptide elevation associated with the development of chronic post-traumatic headache following repetitive blast exposure. *Neurobiology of pain (Cambridge, Mass.)*, 17, 100178.

Ramste M, et al. (2025) Enhancer-targeting CRISPR screens at coronary artery disease loci suggest shared mechanisms of disease risk. *medRxiv : the preprint server for health sciences*.

Xu P, et al. (2025) Defect in hematopoiesis and embryonic lethality at midgestation of Vps13a/Vps13c double knockout mice. *bioRxiv : the preprint server for biology*.

Lu??s IM, et al. (2025) pET28g: A Golden Gate-compatible pET vector for protein expression in *Escherichia coli*, validated by production of functional human ACE2. *PloS one*, 20(7), e0327341.

Holzschuh A, et al. (2025) Rapid multiplexed nanopore amplicon sequencing to distinguish *Plasmodium falciparum* recrudescence from new infection in antimalarial drug trials. *Scientific reports*, 15(1), 36941.

Unitt A, et al. (2025) *Neisseria gonorrhoeae* LIN codes provide a robust, multi-resolution lineage nomenclature. *eLife*, 14.

White BE, et al. (2025) Serial 'deep-sampling' PCR of fragmented DNA reveals the wide range of *Trypanosoma cruzi* burden among chronically infected human, macaque, and canine hosts, and allows accurate monitoring of parasite load following treatment. *eLife*, 14.

Glaser A, et al. (2025) Expansion-assisted selective plane illumination microscopy for nanoscale imaging of centimeter-scale tissues. *eLife*, 12.

Curatolo M, et al. (2025) Patient phenotyping for molecular profiling of neck and low back pain - Study protocol. *Neurobiology of pain (Cambridge, Mass.)*, 18, 100186.

Bahrami A, et al. (2025) Automated 1D Helmholtz coil design for cell biology: Weak magnetic fields alter cytoskeleton dynamics. *PloS one*, 20(8), e0321133.

Punzi S, et al. (2025) Early tolerance and late persistence as alternative drug responses in cancer. *Nature communications*, 16(1), 1291.

Kikawa C, et al. (2025) Near real-time data on the human neutralizing antibody landscape to influenza virus to inform vaccine-strain selection in September 2025. *bioRxiv : the preprint server for biology*.

Pothikamjorn T, et al. (2025) Comparative electron microscopy analysis of internal limiting membrane and epiretinal membrane ultrastructure from vitrectomy surgery: A study

protocol. PloS one, 20(9), e0331248.

Chan LK, et al. (2025) Extrachromosomal circular DNA promotes inflammation and hepatocellular carcinoma development. Science advances, 11(42), eadw0272.

Aho VTE, et al. (2025) A Non-Metaproteomics Researchers' View on Metaproteomics in Microbiome Research. Proteomics, 25(17-18), 28.

Chakraborty D, et al. (2025) Assessment of the Diagnostic Accuracy of the Modified Hodge Test and Modified Carbapenem Inactivation Method for Identifying Carbapenem Resistance Mechanisms in *Klebsiella pneumoniae*: A Whole Genome Sequencing-Based Exploratory Study. Cureus, 17(3), e80312.

Gray CN, et al. (2025) Integrator complex subunit 12 knockout overcomes a transcriptional block to HIV latency reversal. eLife, 13.

Duong TE, et al. (2025) A Multimodal Spatial and Epigenomic Atlas of Human Adult Lung Topography. bioRxiv : the preprint server for biology.

Mitchell E, et al. (2025) Hyposmia in Parkinson's disease; exploring selective odour loss. NPJ Parkinson's disease, 11(1), 67.