

Resource Summary Report

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Samba

RRID:SCR_006557

Type: Tool

Proper Citation

Samba (RRID:SCR_006557)

Resource Information

URL: <http://www.samba.org/>

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Description: Standard Windows interoperability suite of programs for Linux and Unix that provides secure, stable and fast file and print services for all clients using the SMB/CIFS protocol, such as all versions of DOS and Windows, OS/2, Linux and many others. Samba is an important component to seamlessly integrate Linux/Unix Servers and Desktops into Active Directory environments using the winbind daemon.

Abbreviations: Samba

Resource Type: software application, systems interoperability software, software resource

Keywords: interoperability, linux, unix, windows

Availability: GNU General Public License

Resource Name: Samba

Resource ID: SCR_006557

Alternate IDs: nlx_156712

Record Creation Time: 20220129T080236+0000

Record Last Update: 20260728T043235+0000

Ratings and Alerts

No rating or validation information has been found for Samba.

No alerts have been found for Samba.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 123 mentions in open access literature.

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

Sari E, et al. (2026) Gene duplication, horizontal gene transfer, and trait trade-offs drive evolution of postfire resource acquisition in pyrophilous fungi. *Proceedings of the National Academy of Sciences of the United States of America*, 123(1), e2519152123.

Wen H, et al. (2025) TRFill: synergistic use of HiFi and Hi-C sequencing enables accurate assembly of tandem repeats for population-level analysis. *Genome biology*, 26(1), 227.

Wertz AE, et al. (2025) A semisynthetic, multicofactor artificial metalloenzyme retains independent site activity. *Journal of biological inorganic chemistry : JBIC : a publication of the Society of Biological Inorganic Chemistry*, 30(1), 13.

Atkin C, et al. (2025) Provision of medical same day emergency care services within the UK: analysis from the Society for Acute Medicine Benchmarking Audit. *BMJ open*, 15(4), e094580.

Bridgeford EW, et al. (2025) Network Biomarkers of Alzheimer's Disease Risk Derived from Joint Volume and Texture Covariance Patterns in Mouse Models. *bioRxiv : the preprint server for biology*.

Cohen-Rengifo M, et al. (2025) The nasal microbiota of two marine fish species: diversity, community structure, variability, and first insights into the impacts of climate change-related stressors. *FEMS microbiology ecology*, 101(3).

Moon HS, et al. (2025) Smelling the Risk: Early Olfactory Deficits, Brain Networks, and Blood Markers of Alzheimer's Disease Risk in Humanized APOE Mice. *bioRxiv : the preprint server for biology*.

Rousseau C, et al. (2025) A Practical Comparison of Short- and Long-Read Metabarcoding Sequencing: Challenges and Solutions for Plastid Read Removal and Microbial Community Exploration of Seaweed Samples. *Molecular ecology resources*, 25(7), e14129.

Mansour H, et al. (2025) The Duke Mouse Brain Atlas: MRI and light sheet microscopy stereotaxic atlas of the mouse brain. *Science advances*, 11(18), eadq8089.

Paulino S, et al. (2025) Drivers of seasonal dynamics in *Ulva* spp. associated microbiota and surface metabolome: The interplay between environment and host physiology. *Current research in microbial sciences*, 9, 100439.

Zhang H, et al. (2025) Spatially resolved C1QC+ macrophage-CD4+ T cell niche in colorectal cancer microenvironment: implications for immunotherapy response. *Cell*

discovery, 11(1), 60.

Venkadesh S, et al. (2025) A hierarchical framework for cortical and subcortical gray-matter parcellation across rodents, primates, and humans. *bioRxiv : the preprint server for biology*.

Ghiglione JF, et al. (2025) Mission Tara Microplastics: a holistic set of protocols and data resources for the field investigation of plastic pollution along the land-sea continuum in Europe. *Environmental science and pollution research international*, 32(16), 10032.

Wang X, et al. (2025) Review of Methods for Studying Viruses in the Environment and Organisms. *Viruses*, 17(1).

Roberts WR, et al. (2025) Three reference genomes for freshwater diatom ecology and evolution. *Journal of phycology*, 61(2), 267.

Chen CC, et al. (2025) Temperature-dependent shifts in gut microbiota and metabolome of olive flounder (*Paralichthys olivaceus*): implications for cold-water aquaculture expansion and probiotic applications. *Animal microbiome*, 7(1), 49.

Bridgeford EW, et al. (2025) Network biomarkers of Alzheimer's disease risk derived from joint volume and texture covariance patterns in mouse models. *PloS one*, 20(8), e0327118.

Moroni F, et al. (2025) Beyond Microbial Variability: Disclosing the Functional Redundancy of the Core Gut Microbiota of Farmed Gilthead Sea Bream from a Bayesian Network Perspective. *Microorganisms*, 13(1).

Valderrama B, et al. (2025) The South American MicroBiome Archive (saMBA): enriching the microbiome field by studying neglected populations. *Nature communications*, 16(1), 7371.

Choi Y, et al. (2025) All-printed chip-less wearable neuromorphic system for multimodal physicochemical health monitoring. *Nature communications*, 16(1), 5689.