Resource Summary Report

Generated by NIF on Apr 25, 2025

Debian

RRID:SCR_006638
Type: Tool

Proper Citation

Debian (RRID:SCR_006638)

Resource Information

URL: http://www.debian.org

Proper Citation: Debian (RRID:SCR_006638)

Description: Debian is Linux distribution composed of free and open source software, developed by community supported Debian Project, which was established by Ian Murdock on August 16, 1993.Debian comes with over 59000 packages (precompiled software that is bundled up in nice format for easy installation on your machine), package manager (APT), and other utilities that make it possible to manage thousands of packages on thousands of computers as easily as installing single application.

Abbreviations: Debian

Synonyms: Debian - The universal operating system, Debian GNU/Linux

Resource Type: software repository, data or information resource, database, software resource, source code

Keywords: operating system, software package, FASEB list

Funding:

Availability: Free, Freely available

Resource Name: Debian

Resource ID: SCR_006638

Alternate IDs: nlx_151598

License: GNU General Public License

Record Creation Time: 20220129T080237+0000

Record Last Update: 20250425T055543+0000

Ratings and Alerts

No rating or validation information has been found for Debian.

No alerts have been found for Debian.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 47 mentions in open access literature.

Listed below are recent publications. The full list is available at NIF.

Ostermeyer-Fay AG, et al. (2025) The steady-state level of plasma membrane ceramide is regulated by neutral sphingomyelinase 2. Journal of lipid research, 66(1), 100719.

Lyons MA, et al. (2024) Use of multiple pharmacodynamic measures to deconstruct the Nix-TB regimen in a short-course murine model of tuberculosis. Antimicrobial agents and chemotherapy, 68(5), e0101023.

Gómez-Valverde JJ, et al. (2024) Chest X-Ray-Based Telemedicine Platform for Pediatric Tuberculosis Diagnosis in Low-Resource Settings: Development and Validation Study. JMIR pediatrics and parenting, 7, e51743.

Youssef A, et al. (2024) TelePi: an affordable telepathology microscope camera system anyone can build and use. Virchows Archiv : an international journal of pathology, 485(1), 115.

Centanni SW, et al. (2023) PiRATeMC: A highly flexible, scalable, and low-cost system for obtaining high quality video recordings for behavioral neuroscience. Addiction neuroscience, 8.

Greene M, et al. (2023) A simple, highly sensitive, and facile method to quantify ceramide at the plasma membrane. Journal of lipid research, 64(2), 100322.

Lyons MA, et al. (2021) Pretomanid dose selection for pulmonary tuberculosis: An application of multi-objective optimization to dosage regimen design. CPT: pharmacometrics

& systems pharmacology, 10(3), 211.

Park KS, et al. (2021) Cardiac pathologies in mouse loss of imprinting models are due to misexpression of H19 long noncoding RNA. eLife, 10.

Bausch M, et al. (2021) Concept neurons in the human medial temporal lobe flexibly represent abstract relations between concepts. Nature communications, 12(1), 6164.

Viswanathan S, et al. (2020) A response-locking protocol to boost sensitivity for fMRI-based neurochronometry. Human brain mapping, 41(12), 3420.

Florez H, et al. (2020) Online dashboard and data analysis approach for assessing COVID-19 case and death data. F1000Research, 9, 570.

Millán C, et al. (2020) ALIXE: a phase-combination tool for fragment-based molecular replacement. Acta crystallographica. Section D, Structural biology, 76(Pt 3), 209.

Douguet D, et al. (2020) sensaas: Shape-based Alignment by Registration of Colored Point-based Surfaces. Molecular informatics, 39(8), e2000081.

Gruenstaeudl M, et al. (2019) EMBL2checklists: A Python package to facilitate the user-friendly submission of plant and fungal DNA barcoding sequences to ENA. PloS one, 14(1), e0210347.

Arisdakessian C, et al. (2019) DeepImpute: an accurate, fast, and scalable deep neural network method to impute single-cell RNA-seq data. Genome biology, 20(1), 211.

Reber TP, et al. (2019) Representation of abstract semantic knowledge in populations of human single neurons in the medial temporal lobe. PLoS biology, 17(6), e3000290.

Strozzi F, et al. (2019) Scalable Workflows and Reproducible Data Analysis for Genomics. Methods in molecular biology (Clifton, N.J.), 1910, 723.

Jones DR, et al. (2018) SACCHARIS: an automated pipeline to streamline discovery of carbohydrate active enzyme activities within polyspecific families and de novo sequence datasets. Biotechnology for biofuels, 11, 27.

Bak A, et al. (2018) Towards Intelligent Drug Design System: Application of Artificial Dipeptide Receptor Library in QSAR-Oriented Studies. Molecules (Basel, Switzerland), 23(8).

Nastase SA, et al. (2018) Neural Responses to Naturalistic Clips of Behaving Animals in Two Different Task Contexts. Frontiers in neuroscience, 12, 316.