Resource Summary Report

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Launchpad

RRID:SCR_006853
Type: Tool

Proper Citation

Launchpad (RRID:SCR_006853)

Resource Information

URL: https://launchpad.net/

Proper Citation: Launchpad (RRID:SCR_006853)

Description: A software collaboration platform that provides: Bug tracking, Code hosting using Bazaar, Code reviews, Ubuntu package building and hosting, Translations, Mailing lists, Answer tracking and FAQs, and Specification tracking. Launchpad can host your project''s source code using the Bazaar version control system.

Abbreviations: Launchpad

Resource Type: software application, software development tool, software repository, software resource

Keywords: sharing, software, tracking, translation, computer science, web service, source code

Availability: GNU Affero General Public License, v3

Resource Name: Launchpad

Resource ID: SCR_006853

Alternate IDs: nif-0000-10278

Record Creation Time: 20220129T080238+0000

Record Last Update: 20260912T200236+0000

Ratings and Alerts

No rating or validation information has been found for Launchpad.

No alerts have been found for Launchpad.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 27 mentions in open access literature.

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

Pantoja GV, et al. (2026) Bioactive Peptides from Astrocaryum murumuru Residual Biomass: Enzymatic Hydrolysis, Characterization, and Functional Activities. Foods (Basel, Switzerland), 15(11).

Rolland N, et al. (2024) Identification of Food Spoilage Fungi Using MALDI-TOF MS: Spectral Database Development and Application to Species Complex. Journal of fungi (Basel, Switzerland), 10(7).

Guo P, et al. (2021) Identification and Antifungal Susceptibility Analysis of Stephanoascus ciferrii Complex Species Isolated From Patients With Chronic Suppurative Otitis Media. Frontiers in microbiology, 12, 680060.

Grunt TW, et al. (2020) Membrane disruption, but not metabolic rewiring, is the key mechanism of anticancer-action of FASN-inhibitors: a multi-omics analysis in ovarian cancer. Scientific reports, 10(1), 14877.

Kober SL, et al. (2019) Quantification of nitroaromatic explosives in contaminated soil using MALDI-TOF mass spectrometry. Analytical and bioanalytical chemistry, 411(23), 5993.

Ansari JM, et al. (2019) Strain-level diversity of commercial probiotic isolates of Bacillus, Lactobacillus, and Saccharomyces species illustrated by molecular identification and phenotypic profiling. PloS one, 14(3), e0213841.

Mawale R, et al. (2019) Mass spectrometric investigation of amorphous Ga-Sb-Se thin films. Scientific reports, 9(1), 10213.

Kolpikova EP, et al. (2019) Does the Format of Preclass Reading Quizzes Matter? An Evaluation of Traditional and Gamified, Adaptive Preclass Reading Quizzes. CBE life sciences education, 18(4), ar52.

Jo SH, et al. (2019) Structural characterization of phosphoethanolamine-modified lipid A from probiotic Escherichia coli strain Nissle 1917. RSC advances, 9(34), 19762.

Mesureur J, et al. (2018) A MALDI-TOF MS database with broad genus coverage for species-level identification of Brucella. PLoS neglected tropical diseases, 12(10), e0006874.

Rados E, et al. (2018) A laser desorption ionization/matrix-assisted laser desorption ionization target system applicable for three distinct types of instruments (LinTOF/curved field RTOF, LinTOF/RTOF and QqRTOF) with different performance characteristics from

three vendors. Rapid communications in mass spectrometry : RCM, 32(8), 649.

Va?hara P, et al. (2018) Intact Cell Mass Spectrometry as a Quality Control Tool for Revealing Minute Phenotypic Changes of Cultured Human Embryonic Stem Cells. Stem cells translational medicine, 7(1), 109.

Deepaisarn S, et al. (2018) Quantifying biological samples using Linear Poisson Independent Component Analysis for MALDI-ToF mass spectra. Bioinformatics (Oxford, England), 34(6), 1001.

Arjona R, et al. (2018) A PUF- and Biometric-Based Lightweight Hardware Solution to Increase Security at Sensor Nodes. Sensors (Basel, Switzerland), 18(8).

Sim JS, et al. (2018) Lack of the ?1,3-Fucosyltransferase Gene (Osfuct) Affects Anther Development and Pollen Viability in Rice. International journal of molecular sciences, 19(4).

Park HG, et al. (2017) Chemical Structure of the Lipid A component of Pseudomonas sp. strain PAMC 28618 from Thawing Permafrost in Relation to Pathogenicity. Scientific reports, 7(1), 2168.

Wagner R, et al. (2017) Multi-level suppression of receptor-PI3K-mTORC1 by fatty acid synthase inhibitors is crucial for their efficacy against ovarian cancer cells. Oncotarget, 8(7), 11600.

Kadeer A, et al. (2017) Plectin is a novel regulator for apical extrusion of RasV12-transformed cells. Scientific reports, 7, 44328.

Grenfell RC, et al. (2016) Identification of Candida haemulonii Complex Species: Use of ClinProTools(TM) to Overcome Limitations of the Bruker Biotyper(TM), VITEK MS(TM) IVD, and VITEK MS(TM) RUO Databases. Frontiers in microbiology, 7, 940.

Carrick E, et al. (2016) Development of a MALDI MS-based platform for early detection of acute kidney injury. Proteomics. Clinical applications, 10(7), 732.