

Resource Summary Report

Generated by RRID on Aug 6, 2026

LibreOfficeCalc

RRID:SCR_018013

Type: Tool

Proper Citation

LibreOfficeCalc (RRID:SCR_018013)

Resource Information

URL: <https://www.libreoffice.org/>

Proper Citation: LibreOfficeCalc (RRID:SCR_018013)

Description: Open source software application with data analysis tools and spreadsheet templates used to manage, process and visualize data. LibreOffice Calc is spreadsheet component of LibreOffice software package developed by Document Foundation.

Resource Type: data processing software, data management software, software application, data visualization software, data analysis software, software resource

Keywords: Data, analysis, spreadsheet, template, process, visualize

Availability: Free, Available for download, Freely available

Resource Name: LibreOfficeCalc

Resource ID: SCR_018013

Record Creation Time: 20220129T080338+0000

Record Last Update: 20260806T054656+0000

Ratings and Alerts

No rating or validation information has been found for LibreOfficeCalc.

No alerts have been found for LibreOfficeCalc.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 24 mentions in open access literature.

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

Dutkiewicz Z, et al. (2025) Proposal of *Patescibacterium danicum* gen. nov., sp. nov. in the ubiquitous bacterial phylum Patescibacteriota phyl. nov. ISME communications, 5(1), ycae147.

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.

Descarpentrie J, et al. (2024) Protocol for open-source automated universal high-content multiplex fluorescence for RNA in situ analysis. STAR protocols, 5(4), 103508.

Uhlman K, et al. (2024) Routine Pathological Examination of Clinically Presumed Dupuytren Disease: Does It Add Value? Hand (New York, N.Y.), 19(5), 783.

Audebert L, et al. (2024) RNA degradation triggered by decapping is largely independent of initial deadenylation. The EMBO journal.

Karmakar S, et al. (2024) Uniform dipole resonance and suppressed quadrupole resonance for constant transmittivity full phase control plasmonic metasurfaces. Scientific reports, 14(1), 31499.

Bonert M, et al. (2024) Neuroanatomical location of lung cancer brain metastases in 234 patients with a focus on cancer subtyping and biomarkers. PloS one, 19(11), e0314205.

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.

Mendieta D, et al. (2023) Insights into the Genetic Diversity of *Leishmania (Viannia) panamensis* in Panama, Inferred via Multilocus Sequence Typing (MLST). Pathogens (Basel, Switzerland), 12(5).

Le Clercq LS, et al. (2023) Birds of a feather flock together: a dataset for Clock and Adcyap1 genes from migration genetics studies. Scientific data, 10(1), 787.

Dediu D, et al. (2022) The heritability of vocal tract structures estimated from structural MRI in a large cohort of Dutch twins. Human genetics, 141(12), 1905.

Barui S, et al. (2022) Perception without preconception: comparison between the human and machine learner in recognition of tissues from histological sections. Scientific reports, 12(1), 16420.

Zago E, et al. (2022) Early downregulation of hsa-miR-144-3p in serum from drug-naïve Parkinson's disease patients. Scientific reports, 12(1), 1330.

Calvet J, et al. (2022) Biomarker candidates for progression and clinical management of COVID-19 associated pneumonia at time of admission. Scientific reports, 12(1), 640.

Manzini S, et al. (2021) reString: an open-source Python software to perform automatic functional enrichment retrieval, results aggregation and data visualization. *Scientific reports*, 11(1), 23458.

Winkler R, et al. (2021) ProtyQuant: Comparing label-free shotgun proteomics datasets using accumulated peptide probabilities. *Journal of proteomics*, 230, 103985.

Ebert LC, et al. (2021) A review of visualization techniques of post-mortem computed tomography data for forensic death investigations. *International journal of legal medicine*, 135(5), 1855.

Prevosti FJ, et al. (2021) New radiometric ^{40}Ar - ^{39}Ar dates and faunistic analyses refine evolutionary dynamics of Neogene vertebrate assemblages in southern South America. *Scientific reports*, 11(1), 9830.

Phadke KS, et al. (2021) Novel non intrusive continuous use ZeBox technology to trap and kill airborne microbes. *Scientific reports*, 11(1), 22779.

Ksiezopolska E, et al. (2021) Narrow mutational signatures drive acquisition of multidrug resistance in the fungal pathogen *Candida glabrata*. *Current biology : CB*, 31(23), 5314.