

# Resource Summary Report

Generated by [NIF](#) on Sep 12, 2026

## LibreOfficeCalc

RRID:SCR\_018013

Type: Tool

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### Proper Citation

LibreOfficeCalc (RRID:SCR\_018013)

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### 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 analysis software, data management software, data processing software, data visualization software, software application, 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:** 20260912T195856+0000

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### Ratings and Alerts

No rating or validation information has been found for LibreOfficeCalc.

No alerts have been found for LibreOfficeCalc.

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### Data and Source Information

**Source:** [SciCrunch Registry](#)

## Usage and Citation Metrics

We found 33 mentions in open access literature.

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

Krzyżciak P, et al. (2026) Fingernail Onychomycosis: A Laboratory-Based Retrospective Study with Species Profiling and Antifungal Susceptibility of Yeasts. *Journal of clinical medicine*, 15(1).

Jakubczyk D, et al. (2026) Comprehensive Protocol for Handling Human Small Airway Epithelial Cells (HSAECs) to Establish Air-Liquid Interface (ALI) Cultures With TEER-Based Barrier Integrity Assessment. *Bio-protocol*, 16(11), e5699.

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.

Manske F, et al. (2025) Development of a new data management system for the study of the gut microbiome of children who are small for their gestational age. *Computational and structural biotechnology journal*, 27, 221.

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.

Bertin D, et al. (2024) Comparison of the Capacity of Several Machine Learning Tools to Assist Immunofluorescence-Based Detection of Anti-Neutrophil Cytoplasmic Antibodies. *International journal of molecular sciences*, 25(6).

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.

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

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.

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.

Dubey SM, et al. (2023) Relative Membrane Potential Measurements Using DISBAC2(3) Fluorescence in *Arabidopsis thaliana* Primary Roots. *Bio-protocol*, 13(14), e4778.

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.

Chetverikov SF, et al. (2023) Approaches to Sampling for Quality Control of Artificial Intelligence in Biomedical Research. *Sovremennye tekhnologii v meditsine*, 15(2), 19.

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).

Selim M, et al. (2023) The revitalization of endangered heritage buildings: A decision-making framework for investment and determining the highest and best use in Egypt. *F1000Research*, 12, 874.

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.

Kolesov DE, et al. (2022) Antigenic properties of the SARS-CoV-2 nucleoprotein are altered by the RNA admixture. *PeerJ*, 10, e12751.

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.

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.