

Resource Summary Report

Generated by PRECISE-TBI on Sep 7, 2026

streamlit

RRID:SCR_024354

Type: Tool

Proper Citation

streamlit (RRID:SCR_024354)

Resource Information

URL: <https://github.com/streamlit/streamlit>

Proper Citation: streamlit (RRID:SCR_024354)

Description: Software tool to turn data scripts into shareable web apps in minutes. Faster way to build and share data apps.

Resource Type: software application, software resource

Keywords: turn data scripts into shareable web apps, build and share data apps,

Availability: Free, Available for download, Freely available,

Resource Name: streamlit

Resource ID: SCR_024354

Old URLs: <https://sources.debian.org/src/streamlit/>

License: Apache-2.0 license

Record Creation Time: 20230830T050217+0000

Record Last Update: 20260905T133317+0000

Ratings and Alerts

No rating or validation information has been found for streamlit.

No alerts have been found for streamlit.

Data and Source Information

Usage and Citation Metrics

We found 9 mentions in open access literature.

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

Zabetian Z, et al. (2025) Protocol for deep-learning-driven cell type label transfer in single-cell RNA sequencing data. STAR protocols, 6(2), 103768.

Saygili ES, et al. (2025) Machine Learning-Based Survival Prediction Tool for Adrenocortical Carcinoma. The Journal of clinical endocrinology and metabolism, 110(10), e3185.

Brown C, et al. (2025) A proteome-wide quantitative platform for nanoscale spatially resolved extraction of membrane proteins into native nanodiscs. Nature methods, 22(2), 412.

Wang SQ, et al. (2025) Development and validation of web-based, interpretable predictive models for sepsis and mortality in extensive burns. Frontiers in cellular and infection microbiology, 15, 1586087.

Radtke M, et al. (2024) altAFplotter: a web app for reliable UPD detection in NGS diagnostics. BMC bioinformatics, 25(1), 299.

Brown C, et al. (2024) A proteome-wide quantitative platform for nanoscale spatially resolved extraction of membrane proteins into native nanodiscs. bioRxiv : the preprint server for biology.

Laprade W, et al. (2023) Machine-learning guided Venom Induced Dermonecrosis Analysis tool: VIDAL. Scientific reports, 13(1), 21662.

Nápoles-Duarte JM, et al. (2022) Stmol: A component for building interactive molecular visualizations within streamlit web-applications. Frontiers in molecular biosciences, 9, 990846.

Shah W, et al. () Open-source pre-clinical image segmentation: mouse cardiac magnetic resonance imaging datasets with a deep learning segmentation framework. Journal of cardiovascular magnetic resonance : official journal of the Society for Cardiovascular Magnetic Resonance, 28(1), 102706.