

Resource Summary Report

Generated by [RRID](#) on Aug 2, 2026

Stack Overflow

RRID:SCR_011984

Type: Tool

Proper Citation

Stack Overflow (RRID:SCR_011984)

Resource Information

URL: <http://stackoverflow.com/>

Proper Citation: Stack Overflow (RRID:SCR_011984)

Description: A question and answer site for professional and enthusiast programmers.

Abbreviations: Stack Overflow

Resource Type: portal, data or information resource, community building portal

Keywords: programmer

Availability: Free

Resource Name: Stack Overflow

Resource ID: SCR_011984

Alternate IDs: OMICS_01711

Record Creation Time: 20220129T080307+0000

Record Last Update: 20260801T190427+0000

Ratings and Alerts

No rating or validation information has been found for Stack Overflow.

No alerts have been found for Stack Overflow.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 20 mentions in open access literature.

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

Elloumi F, et al. (2025) Protocol to set up the CellMinerCDB pharmacogenomics analysis web application with custom cell line data. STAR protocols, 6(4), 104076.

Mukherjee EM, et al. (2025) Immune Checkpoint Inhibitors as Independent and Synergistic Drivers of SJS/TEN: An Analysis of FAERS. medRxiv : the preprint server for health sciences.

Idkowiak J, et al. (2025) Best practices and tools in R and Python for statistical processing and visualization of lipidomics and metabolomics data. Nature communications, 16(1), 8714.

Mukherjee EM, et al. (2025) Seasonality and Trends in Stevens-Johnson Syndrome/Toxic Epidermal Necrolysis Before and During the COVID-19 Pandemic: A Pharmacovigilance Study. medRxiv : the preprint server for health sciences.

Nguyen H, et al. (2024) Intelligent search system for resume and labor law. PeerJ. Computer science, 10, e1786.

Hoffman AM, et al. (2024) Ten simple rules for teaching an introduction to R. PLoS computational biology, 20(5), e1012018.

Wang Q, et al. (2024) Deep active learning for multi label text classification. Scientific reports, 14(1), 28246.

Dasgupta S, et al. (2023) Protocol for evaluating mechanistic pathways associated with HIV acquisition via nested Least Absolute Shrinkage and Selective Operator analysis. STAR protocols, 4(4), 102628.

de Carvalho VDH, et al. (2022) Mining Public Opinions on COVID-19 Vaccination: A Temporal Analysis to Support Combating Misinformation. Tropical medicine and infectious disease, 7(10).

Oellermann M, et al. (2022) Open Hardware in Science: The Benefits of Open Electronics. Integrative and comparative biology, 62(4), 1061.

Islam S, et al. (2021) Ten simple rules for navigating the computational aspect of an interdisciplinary PhD. PLoS computational biology, 17(2), e1008554.

Kamiński A, et al. (2021) An empirical study of Q&A websites for game developers. Empirical software engineering, 26(6), 115.

Stefaniak F, et al. (2021) AnnapuRNA: A scoring function for predicting RNA-small molecule binding poses. PLoS computational biology, 17(2), e1008309.

Moutidis I, et al. (2021) Community evolution on Stack Overflow. PloS one, 16(6), e0253010.

Hammad M, et al. (2021) Clone-advisor: recommending code tokens and clone methods with deep learning and information retrieval. PeerJ. Computer science, 7, e737.

Gujral H, et al. (2020) Utilization of Time Series Tools in Life-sciences and Neuroscience. Neuroscience insights, 15, 2633105520963045.

Ewers-Saucedo C, et al. (2019) Testing adaptive hypotheses on the evolution of larval life history in acorn and stalked barnacles. Ecology and evolution, 9(19), 11434.

Herodotou C, et al. (2018) Designing citizen science tools for learning: lessons learnt from the iterative development of nQuire. Research and practice in technology enhanced learning, 13(1), 4.

Carson MA, et al. (2016) Approaches to R education in Canadian universities. F1000Research, 5, 2802.

, et al. (2013) On Crowd-verification of Biological Networks. Bioinformatics and biology insights, 7, 307.