

Resource Summary Report

Generated by PRECISE-TBI on Aug 1, 2026

ObservableHQ

RRID:SCR_016173

Type: Tool

Proper Citation

ObservableHQ (RRID:SCR_016173)

Resource Information

URL: <https://beta.observablehq.com/>

Proper Citation: ObservableHQ (RRID:SCR_016173)

Description: Web application for a code and text writing environment. It uses javascript and can be used to produce executable papers.

Resource Type: software resource, data access protocol, web application, application programming interface

Keywords: code, text, environment, writing, paper, research, journal, execute, programming, javascript

Availability: Open source, Freely available, Account required, Account required for full access

Resource Name: ObservableHQ

Resource ID: SCR_016173

License URLs: <https://beta.observablehq.com/terms-of-service>

Record Creation Time: 20220129T080329+0000

Record Last Update: 20260801T042756+0000

Ratings and Alerts

No rating or validation information has been found for ObservableHQ.

No alerts have been found for ObservableHQ.

Data and Source Information

Usage and Citation Metrics

We found 7 mentions in open access literature.

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

Woitowich NC, et al. (2024) Analysis of NIH K99/R00 awards and the career progression of awardees. eLife, 12.

Woitowich NC, et al. (2023) Analysis of NIH K99/R00 Awards and the Career Progression of Awardees. bioRxiv : the preprint server for biology.

Martin DP, et al. (2022) Selection Analysis Identifies Clusters of Unusual Mutational Changes in Omicron Lineage BA.1 That Likely Impact Spike Function. Molecular biology and evolution, 39(4).

Martin DP, et al. (2022) Selection analysis identifies unusual clustered mutational changes in Omicron lineage BA.1 that likely impact Spike function. bioRxiv : the preprint server for biology.

Martin DP, et al. (2021) The emergence and ongoing convergent evolution of the N501Y lineages coincides with a major global shift in the SARS-CoV-2 selective landscape. medRxiv : the preprint server for health sciences.

Ortega DR, et al. (2020) Repurposing a chemosensory macromolecular machine. Nature communications, 11(1), 2041.

Torre D, et al. (2018) BioJupies: Automated Generation of Interactive Notebooks for RNA-Seq Data Analysis in the Cloud. Cell systems, 7(5), 556.