

Statistics for ErnakovichLab/metagenomic_methods_ernakovich_lab

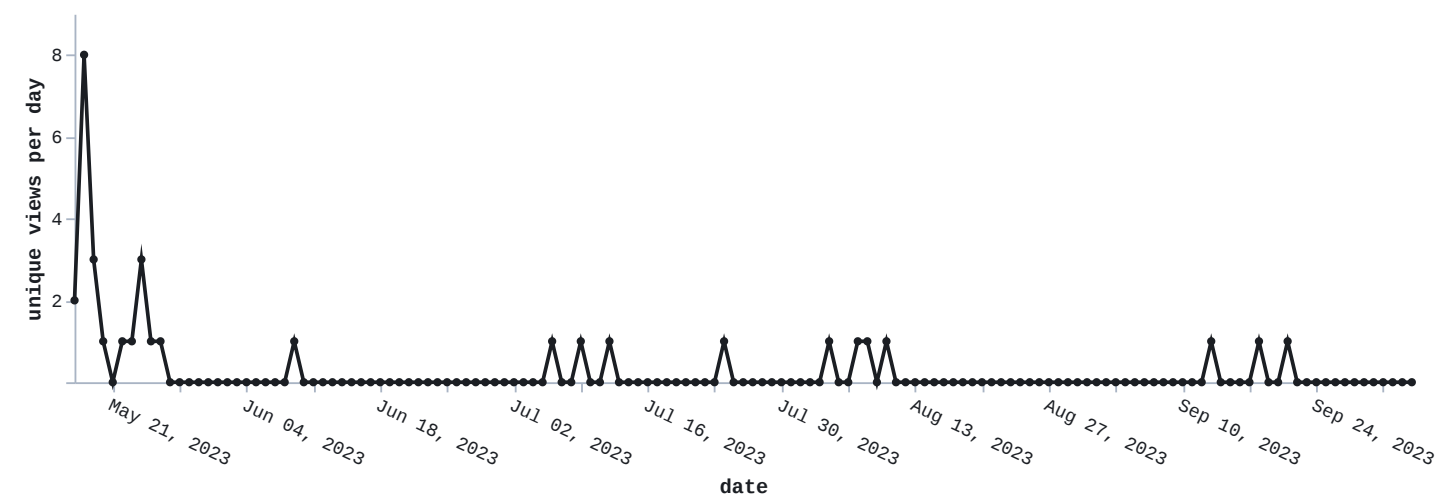
Generated for [ErnakovichLab/metagenomic_methods_ernakovich_lab](#) with [jgehrcke/github-repo-stats](#) at 2023-10-05 23:07 UTC.

Table of contents:

- [Views](#)
- [Clones](#)
- [Stargazers](#)
- [Forks](#)
- [Top referrers and paths](#)

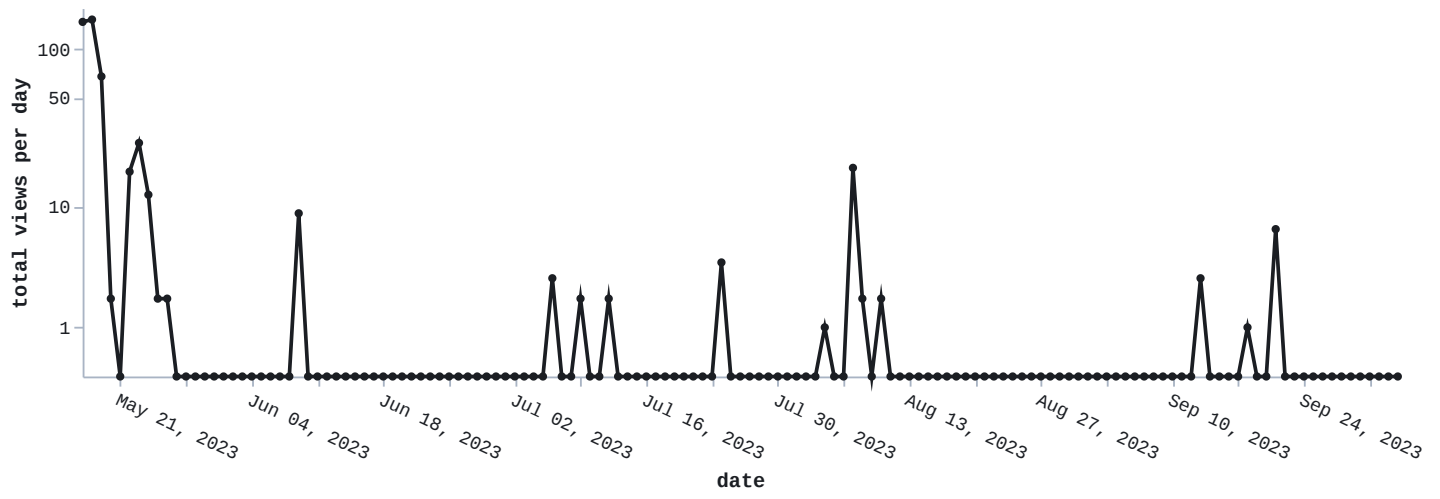
Views

Unique visitors



Cumulative: 33

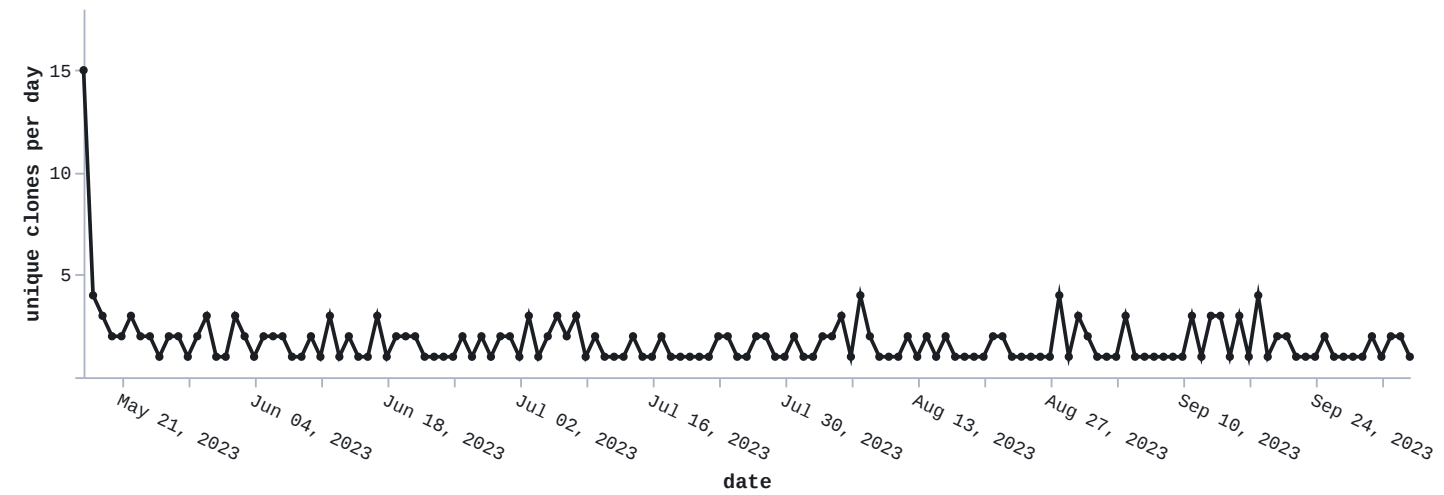
Total views



Cumulative: 484

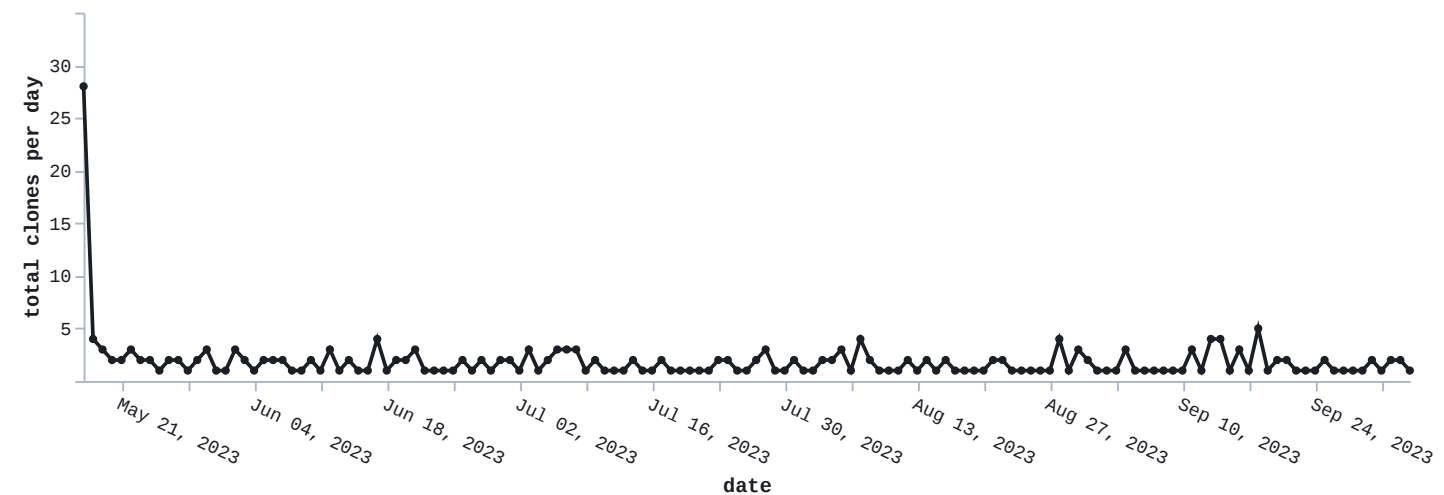
Clones

Unique cloners



Cumulative: 244

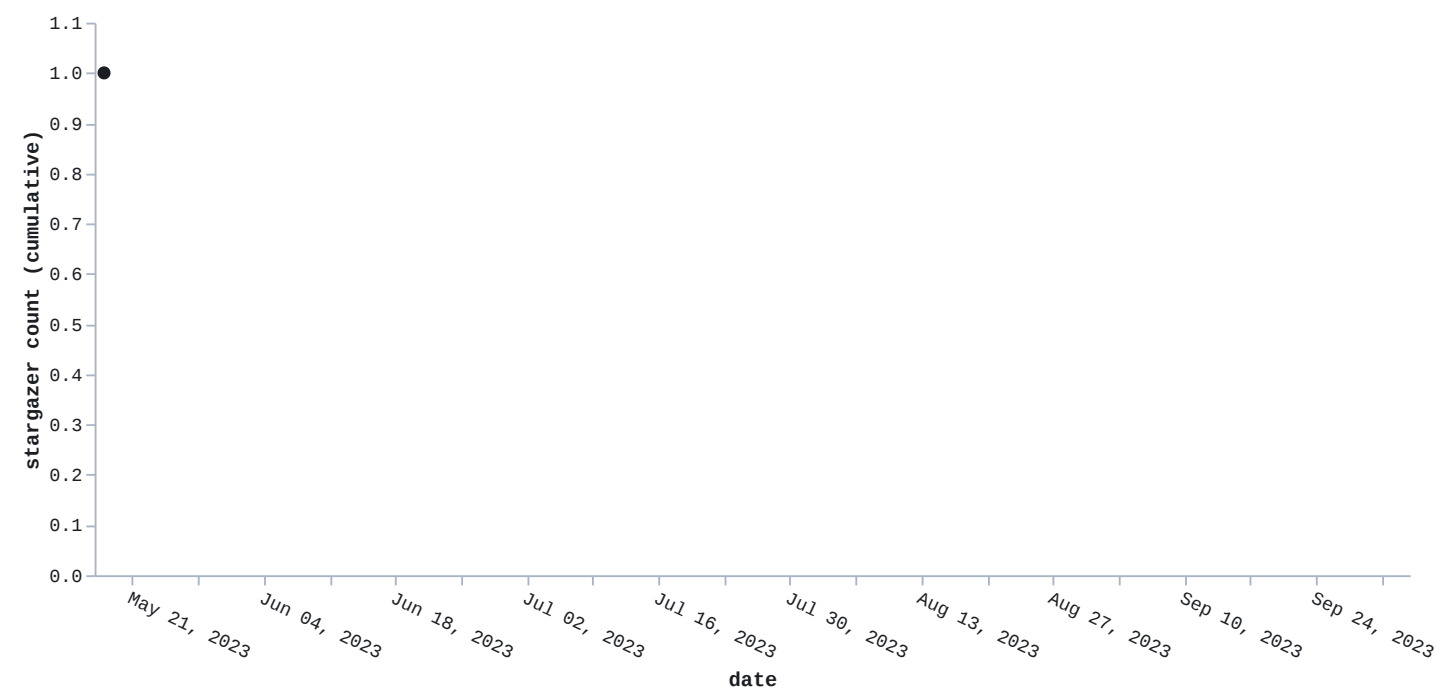
Total clones



Cumulative: 264

Stargazers

Each data point corresponds to at least one stargazer event. The time resolution is one day.



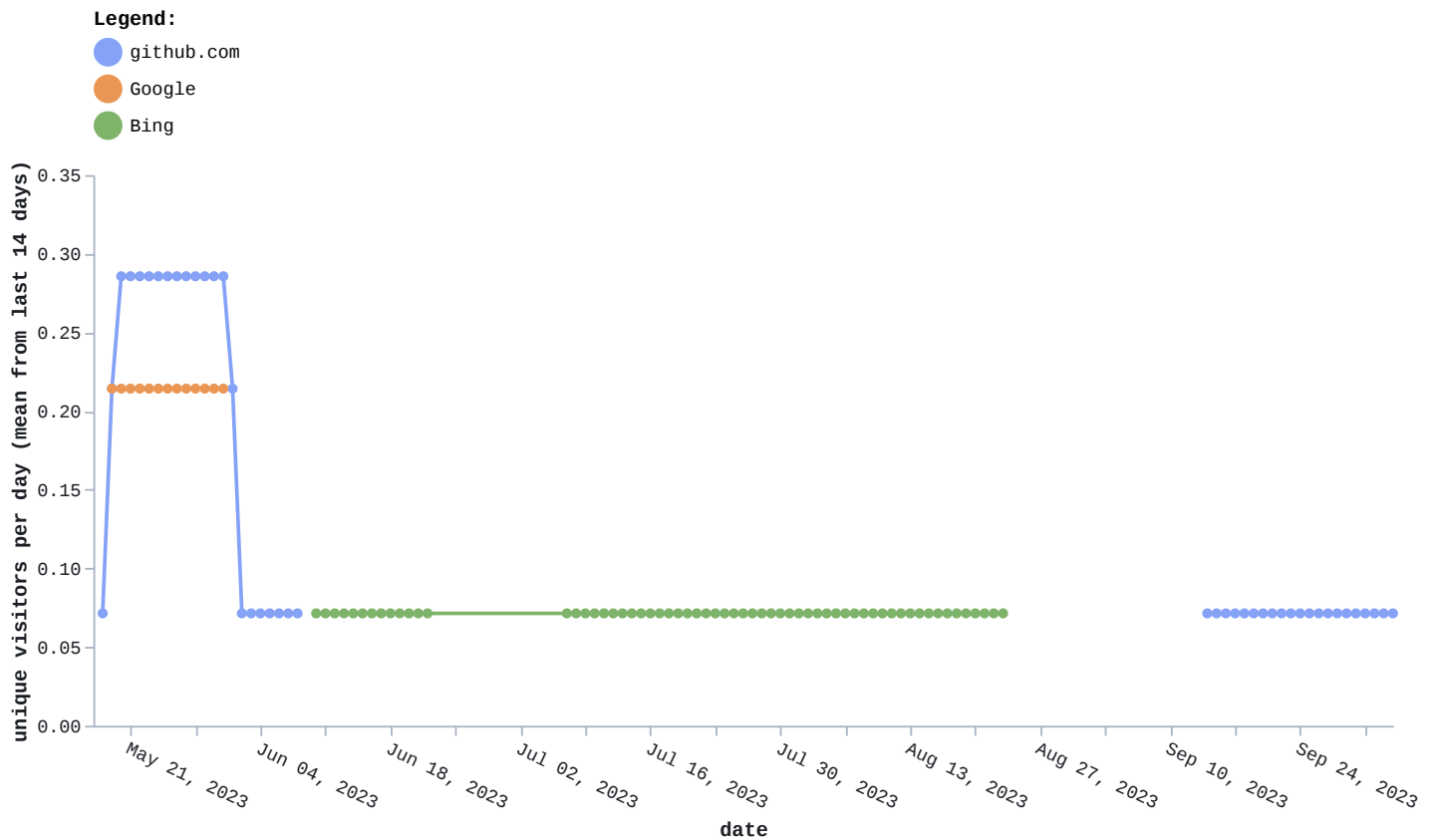
Forks

This repository has no forks yet.

Top referrers and paths

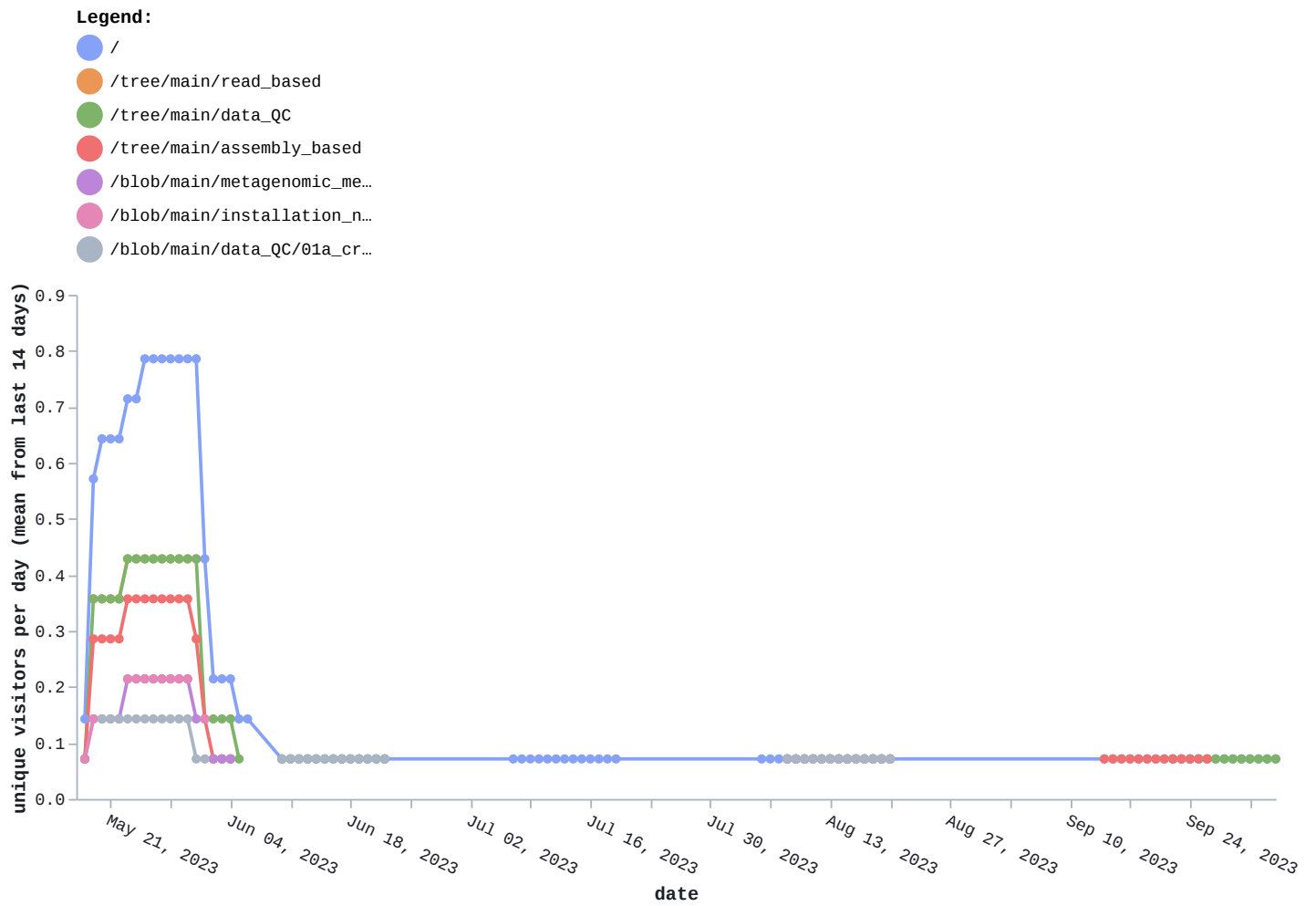
Note: Each data point in the plots shown below is influenced by the 14 days leading up to it. Each data point is the arithmetic mean of the “unique visitors per day” metric, built from a time window of 14 days width, and plotted at the right edge of that very time window. That is, these plots respond slowly to change (narrow peaks are smoothed out).

Top referrers



Top 15 referrers: 01: github.com , 02: Google , 03: Bing

Top paths



Top 15 paths: 01: / , 02: /tree/main/read_based , 03: /tree/main/data_QC , 04: /tree/main/assembly_based , 05: /blob/main/metagenomic_methods_ernakovich_lab.yml , 06: /blob/main/installation_notes.txt , 07: /blob/main/data_QC/01a_create_fastqc_commands.sh , 08: /blob/main/data_QC/01_run_fastqc_commands.slurm , 09: /tree/main , 10: /tree/main/.github/workflows , 11: /blob/main/assembly_based/readme.md , 12: /blob/main/README.md , 13: /blob/main/data_QC/multiqc_config.yaml , 14: /blame/main/data_QC/03a_create_cutadapt_transposase_commands.sh , 15: /blob/main/data_QC/README.md