# Statistics for ErnakovichLab/metagenomic\_methods\_ernakovich lab

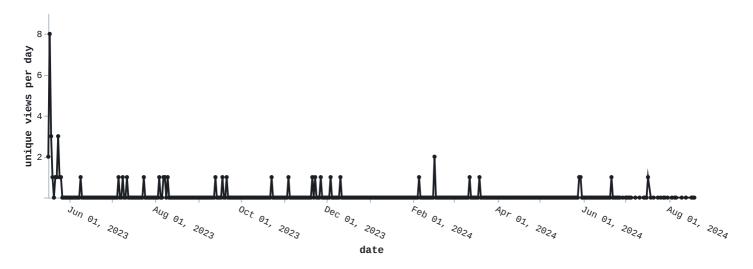
Generated for ErnakovichLab/metagenomic\_methods\_ernakovich\_lab with jgehrcke/github-repo-stats at 2024-08-19 23:10 UTC.

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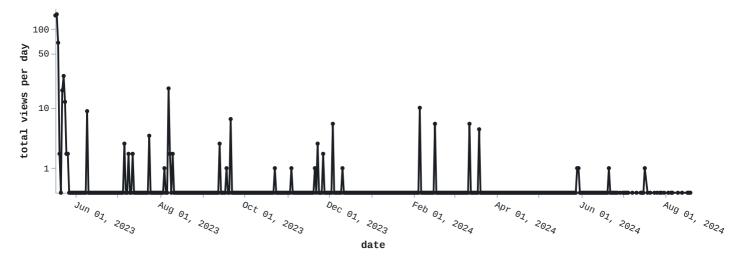
### **Views**

#### **Unique visitors**



Cumulative: 49

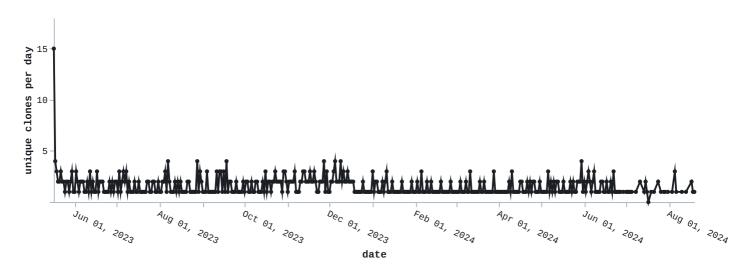
**Total views** 



Cumulative: 530

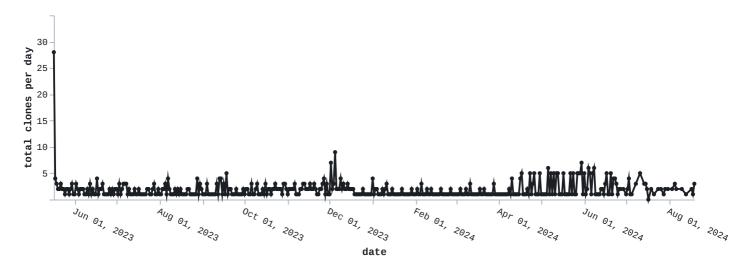
### **Clones**

#### **Unique cloners**



Cumulative: 684

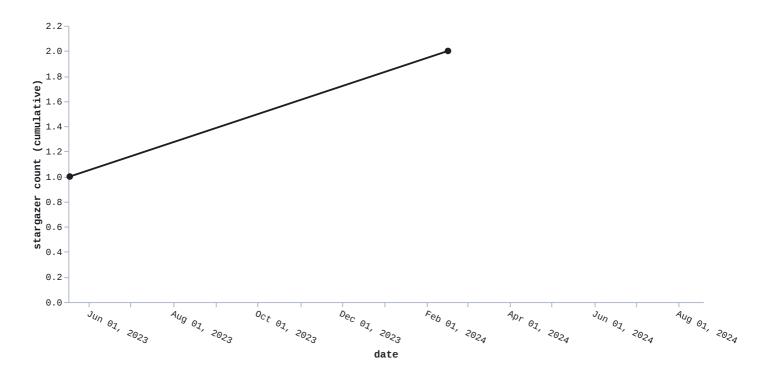
#### **Total clones**



Cumulative: 824

## **Stargazers**

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



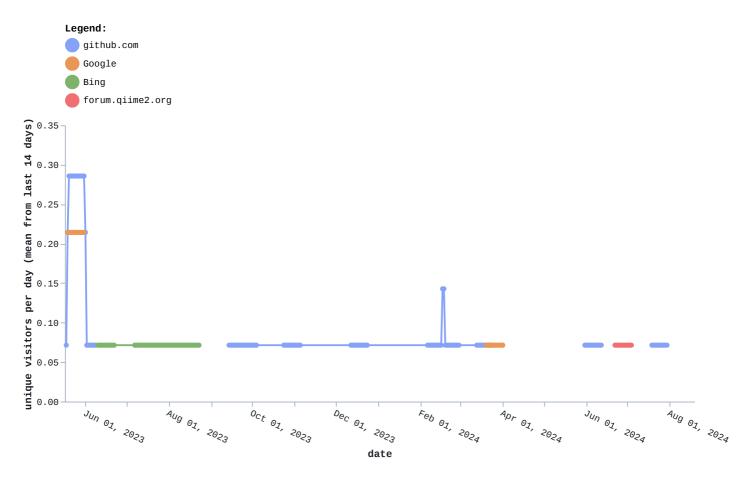
### **Forks**

This repository has no forks yet.

### **Top referrers and paths**

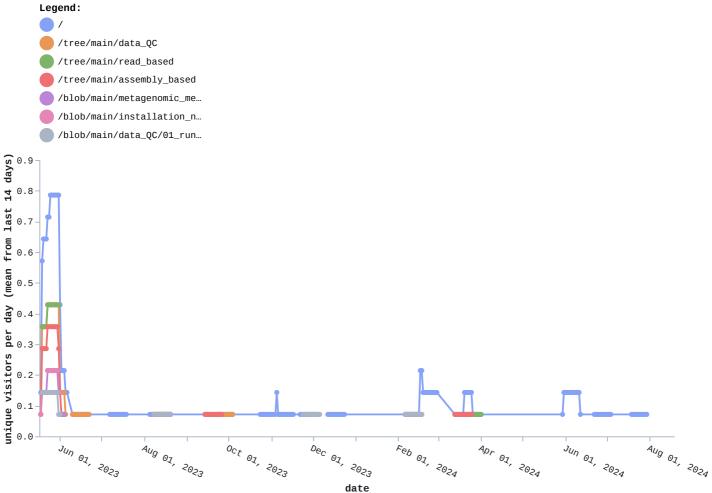
Note: Each data point in the plots shown below is influenced by the 14 days leading up to it. Each data point is the arithmetic mean of the "unique visitors per day" metric, built from a time window of 14 days width, and plotted at the right edge of that very time window. That is, these plots respond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

#### **Top referrers**



Top 15 referrers: 01: github.com, 02: Google, 03: Bing, 04: forum.qiime2.org

#### Top paths



Top 15 paths: 01: /, 02: /tree/main/data\_QC , 03: /tree/main/read\_based , 04: /tree/main/assembly\_based , 05: /blob/main/metagenomic\_methods\_ernakovich\_lab.yml , 06: /blob/main/installation\_notes.txt , 07: /blob/main/data\_QC/01\_run\_fastqc\_commands.slurm , 08: /tree/main/.github/workflows , 09: /blob/main/assembly\_based/readme.md , 10: /blob/main/data\_QC/01a\_create\_fastqc\_commands.sh , 11: /tree/main , 12: /blob/main/data\_QC/README.md , 13: /blob/main/README.md , 14: /blob/main/read\_based/readme.md , 15: /blob/main/data\_QC/multiqc\_config.yaml