

# Statistics for ClaveLab/genome-assembly

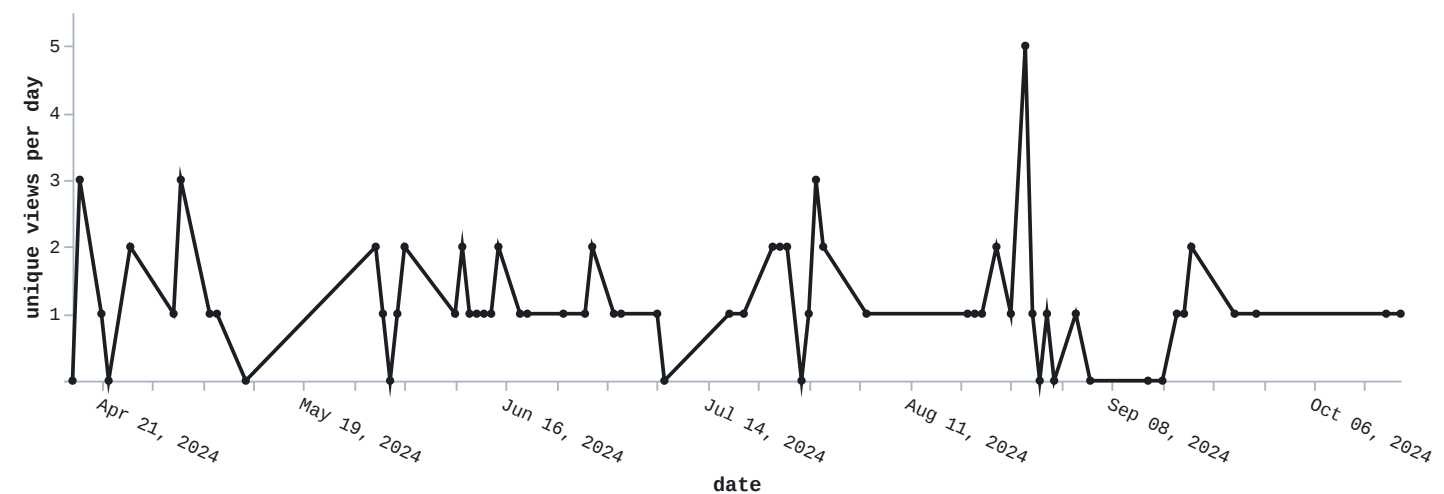
Generated for [ClaveLab/genome-assembly](#) with [jgehrcke/github-repo-stats](#) at 2024-10-20 23:14 UTC.

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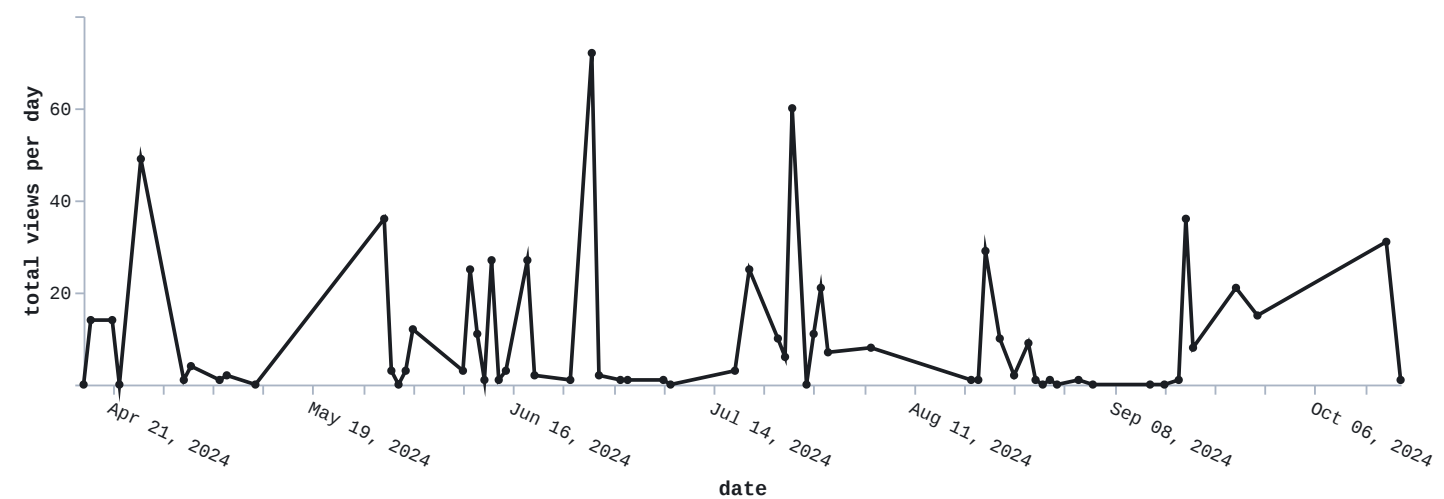
## Views

### Unique visitors



Cumulative: 73

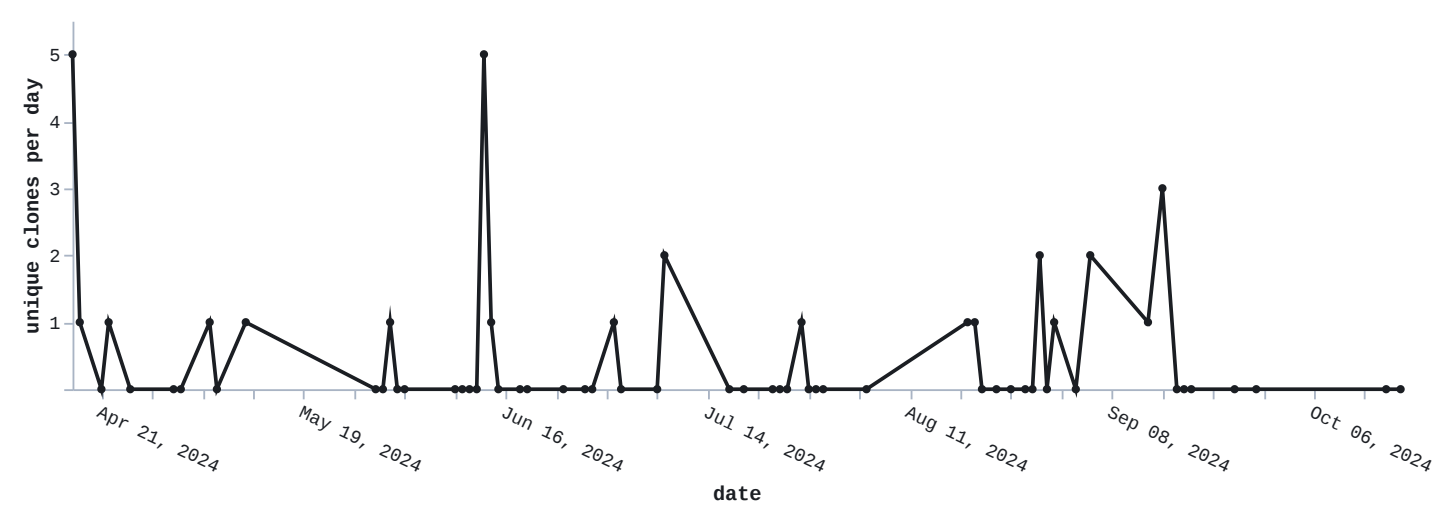
### Total views



Cumulative: 636

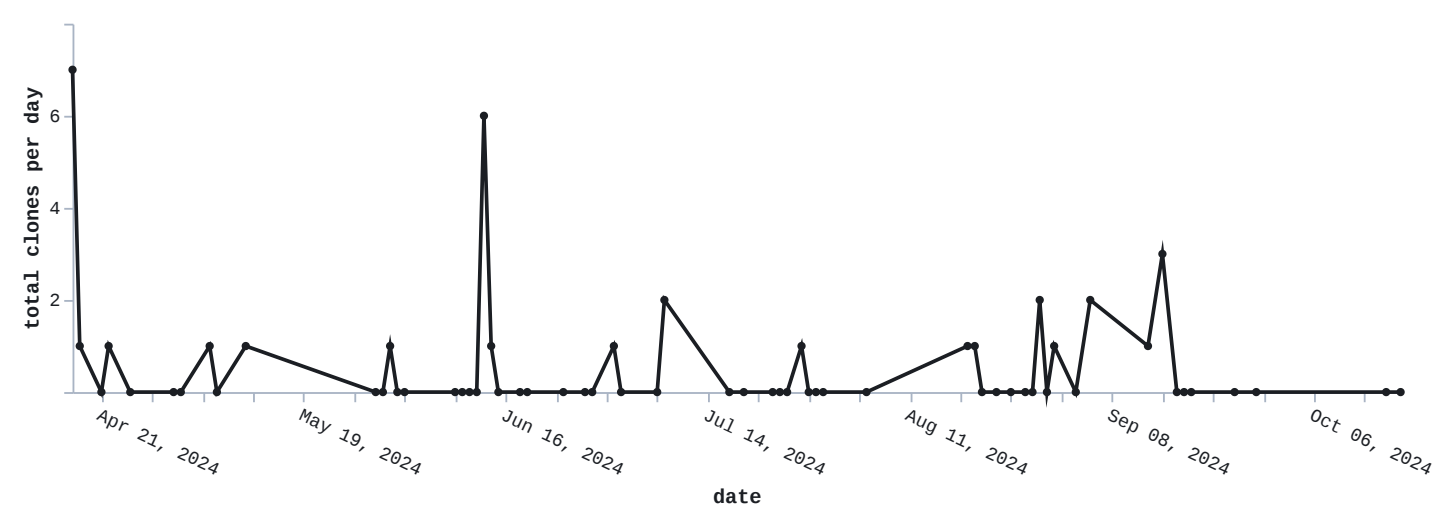
# Clones

## Unique cloners



Cumulative: 31

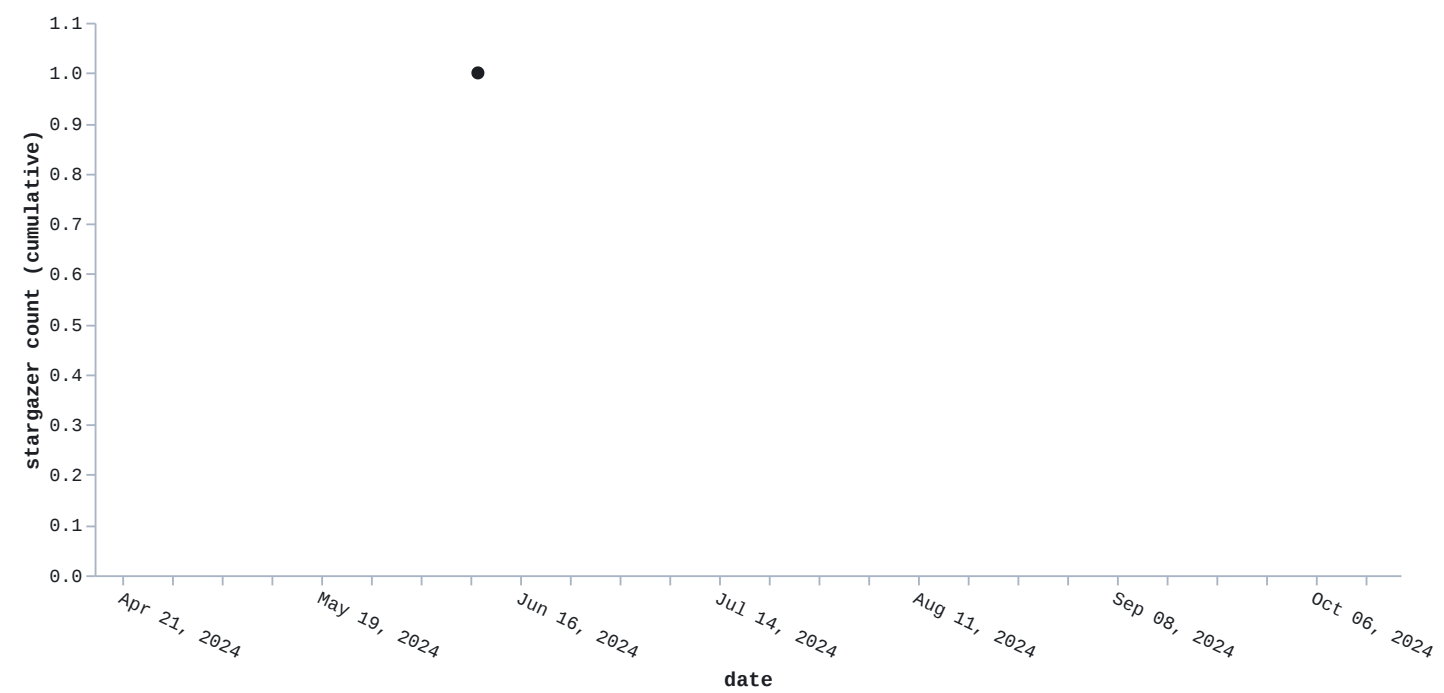
## Total clones



Cumulative: 34

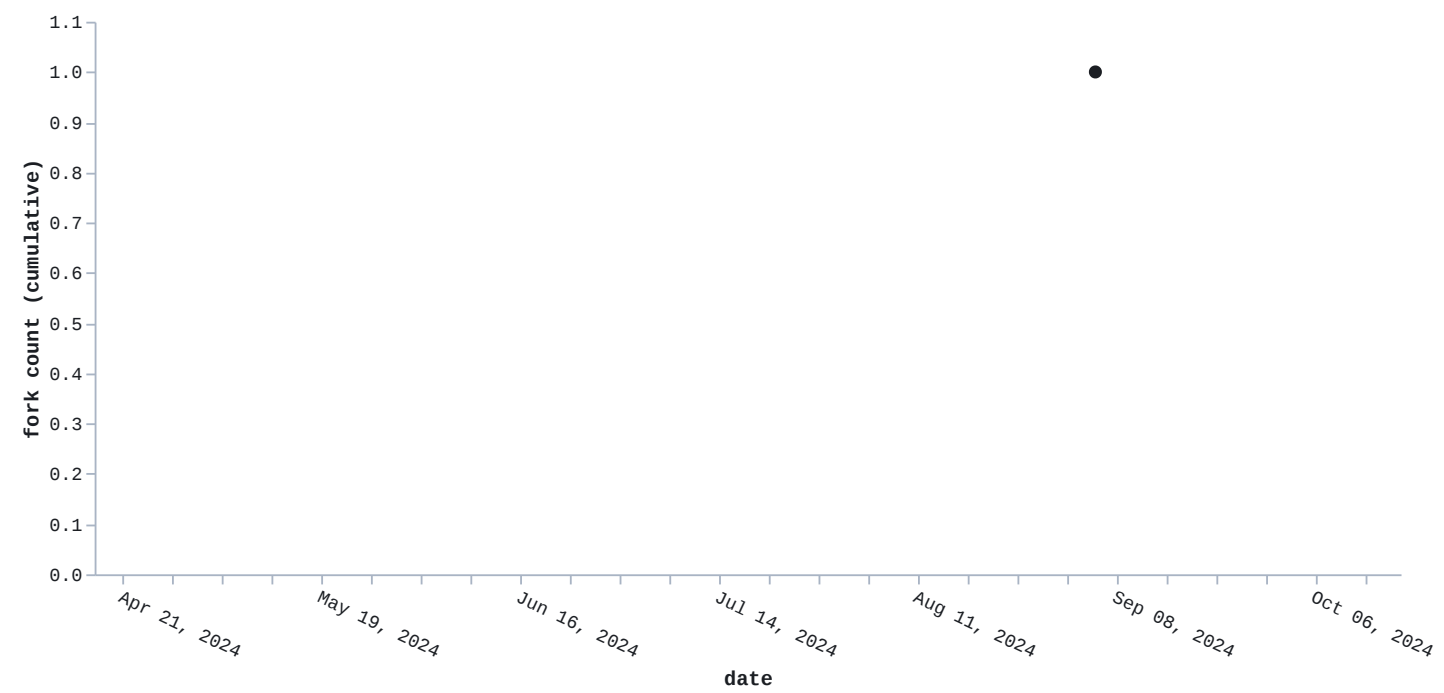
# Stargazers

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



# Forks

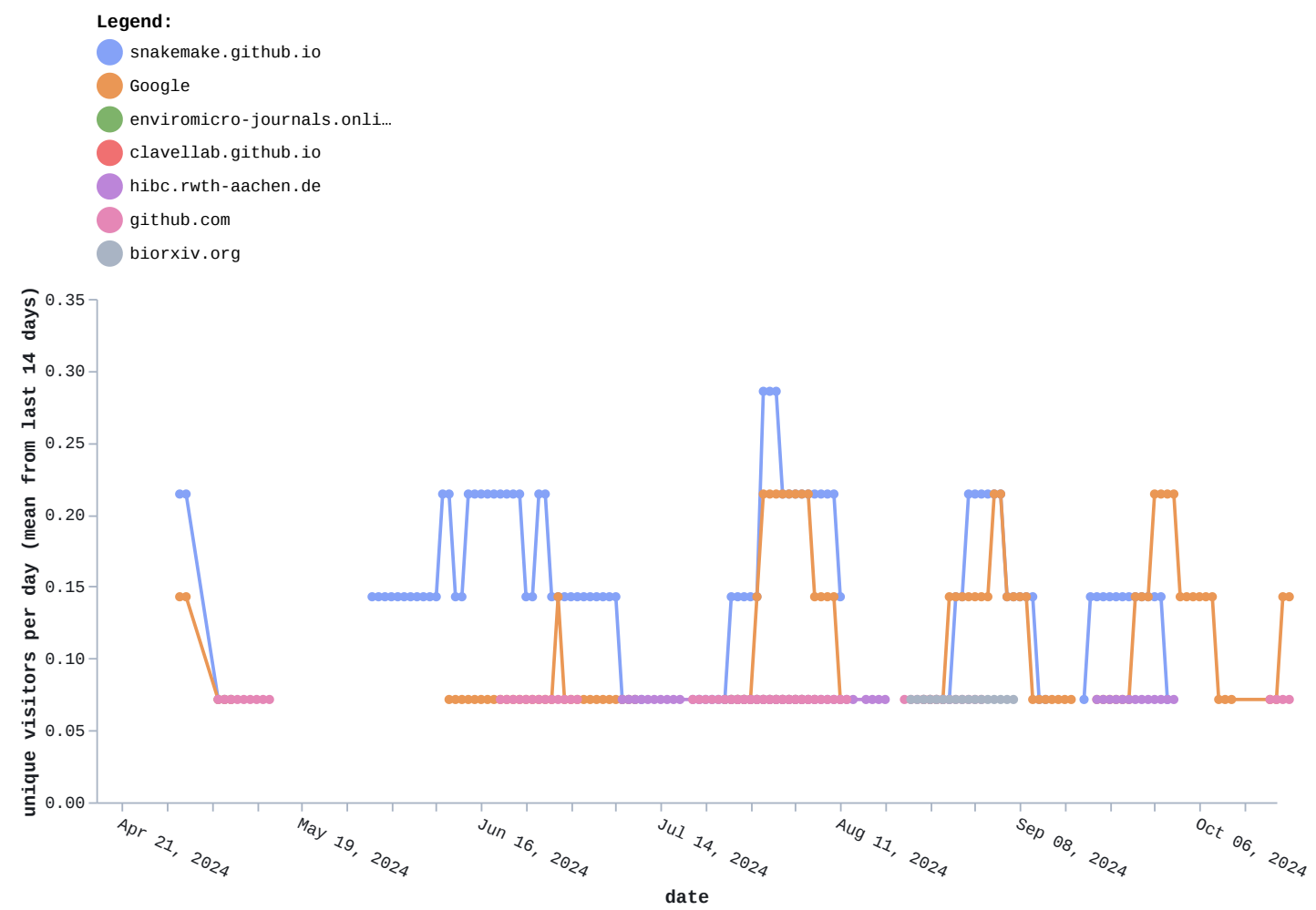
Each data point corresponds to at least one fork event. The time resolution is one day.



# Top referrers and paths

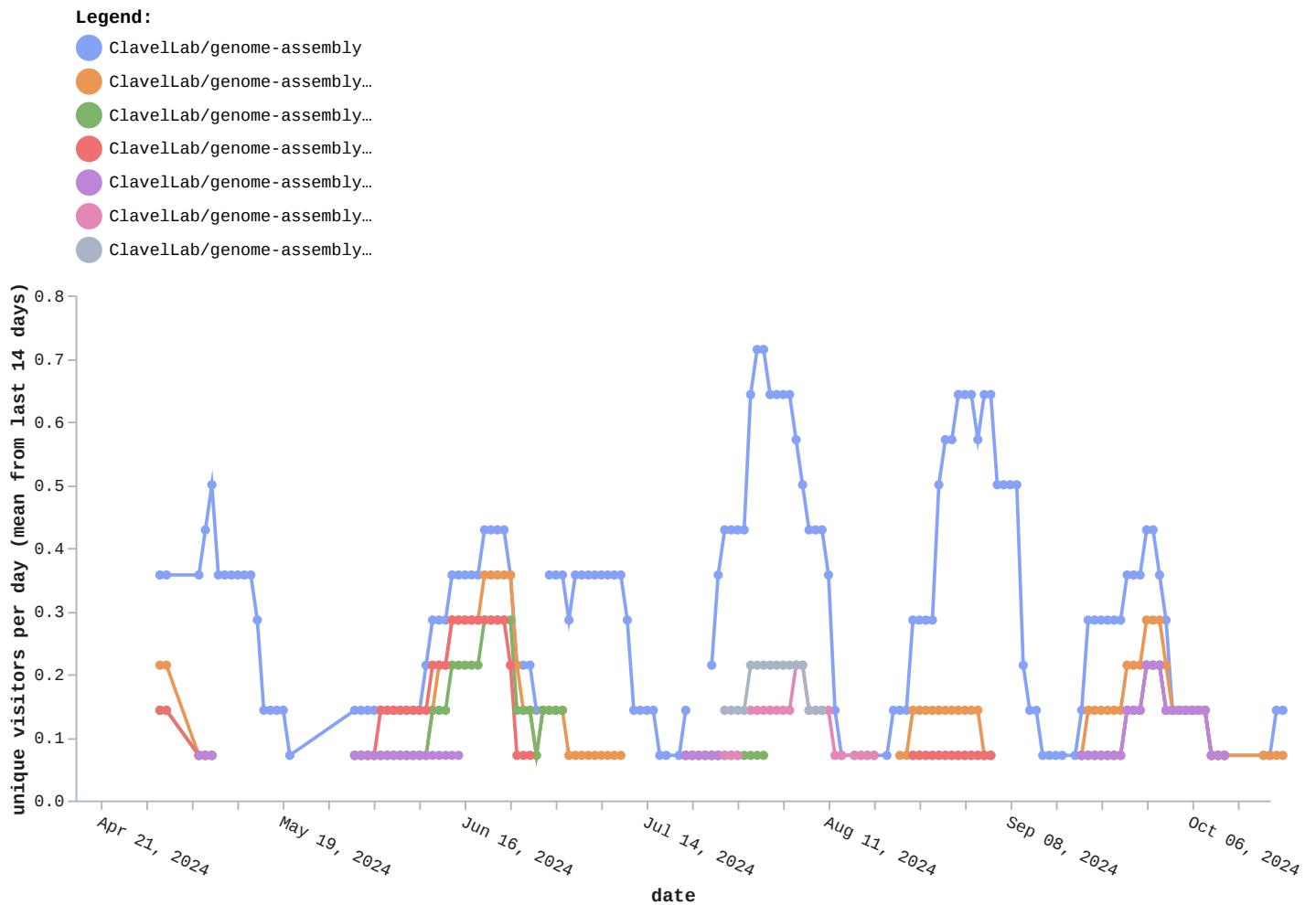
Note: Each data point in the plots shown below is influenced by the 14 days leading up to it. Each data point is the arithmetic mean of the “unique visitors per day” metric, built from a time window of 14 days width, and plotted at the right edge of that very time window. That is, these plots respond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

## Top referrers



Top 15 referrers: 01: snakemake.github.io , 02: Google , 03: enviromicro-journals.onlinelibrary.wiley.com , 04: clavellab.github.io , 05: hibern.rwth-aachen.de , 06: github.com , 07: biorexiv.org

## Top paths



Top 15 paths: 01: ClaveLLab/genome-assembly , 02: ClaveLLab/genome-assembly/tree/main/workflow , 03: ClaveLLab/genome-assembly/tree/main/workflow/rules , 04: ClaveLLab/genome-assembly/blob/main/workflow/rules/assembly.smk , 05: ClaveLLab/genome-assembly/blob/main/workflow/rules/filtering.smk , 06: ClaveLLab/genome-assembly/actions , 07: ClaveLLab/genome-assembly/issues , 08: ClaveLLab/genome-assembly/blob/main/workflow/rules/quality\_check.smk , 09: ClaveLLab/genome-assembly/pulls , 10: ClaveLLab/genome-assembly/blob/main/config/README.md , 11: ClaveLLab/genome-assembly/tree/main/resources/demo , 12: ClaveLLab/genome-assembly/pulse , 13: ClaveLLab/genome-assembly/tree/main/workflow/schemas , 14: ClaveLLab/genome-assembly/blob/main/workflow/Snakefile , 15: ClaveLLab/genome-assembly/tree/main/workflow/envs