

# Statistics for jodithe/Sex\_differences\_genetics\_depression

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Generated for [jodithe/Sex\\_differences\\_genetics\\_depression](#) with [jgehrcke/github-repo-stats](#) at 2026-02-23 23:24 UTC.

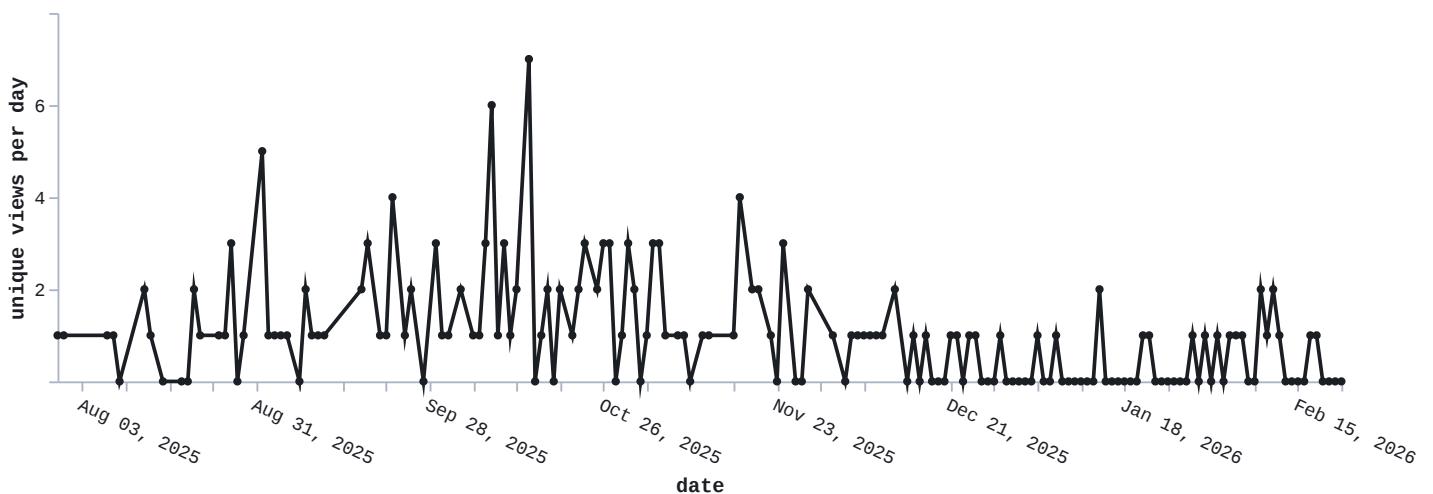
Table of contents:

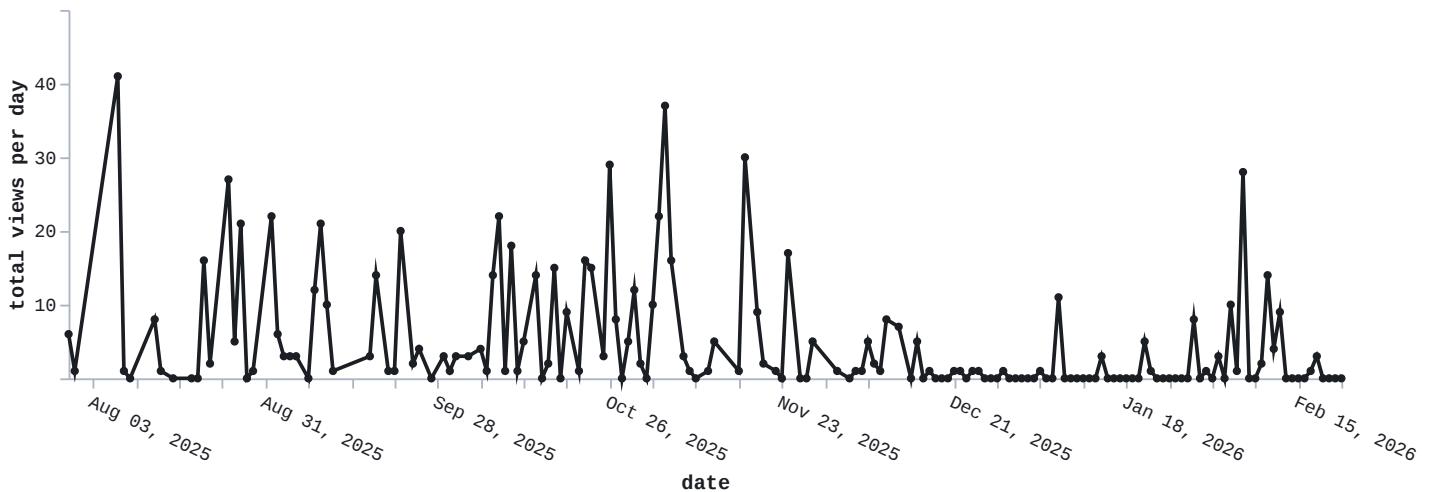
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## Views

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### Unique visitors

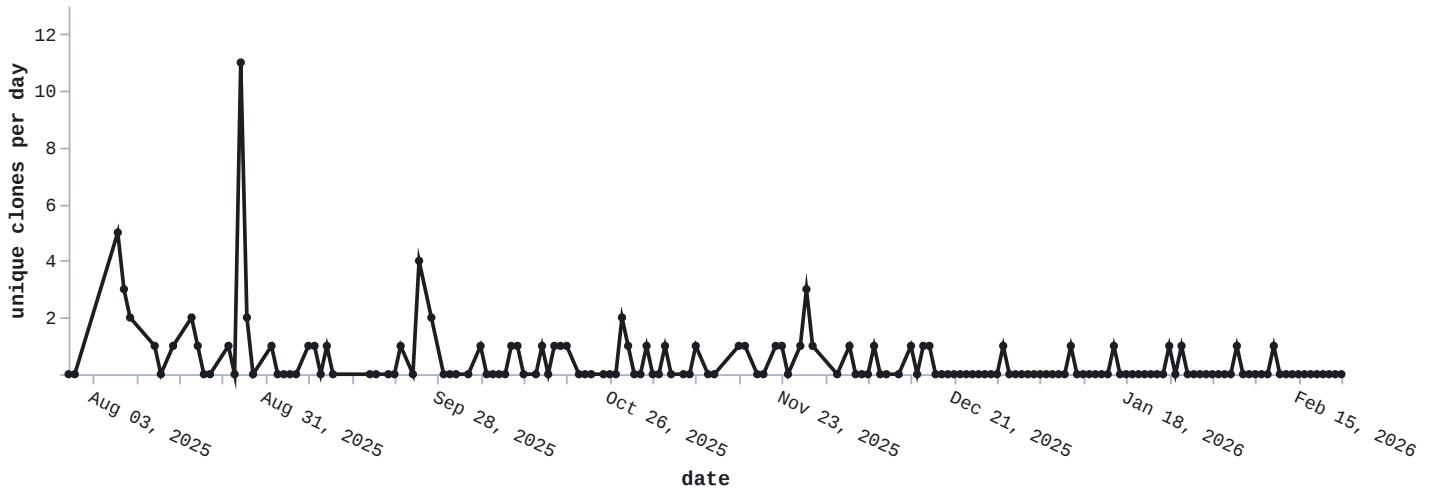




Cumulative: 765

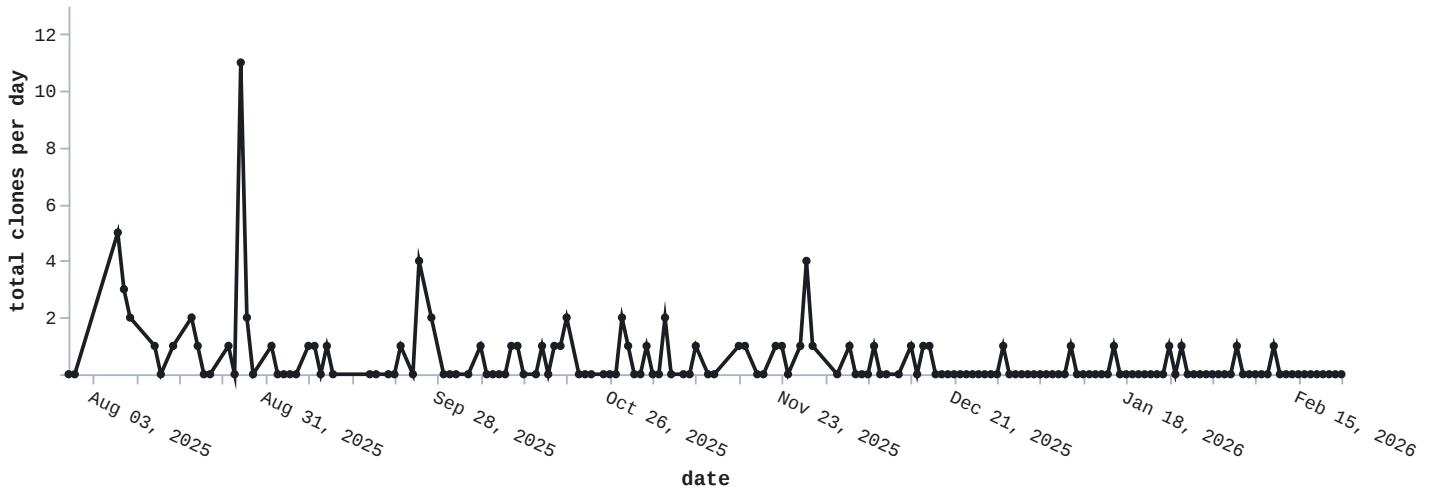
# Clones

## Unique cloners



Cumulative: 74

## Total clones



Cumulative: 77

## **Stargazers**

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This repository has no stars yet.

## **Forks**

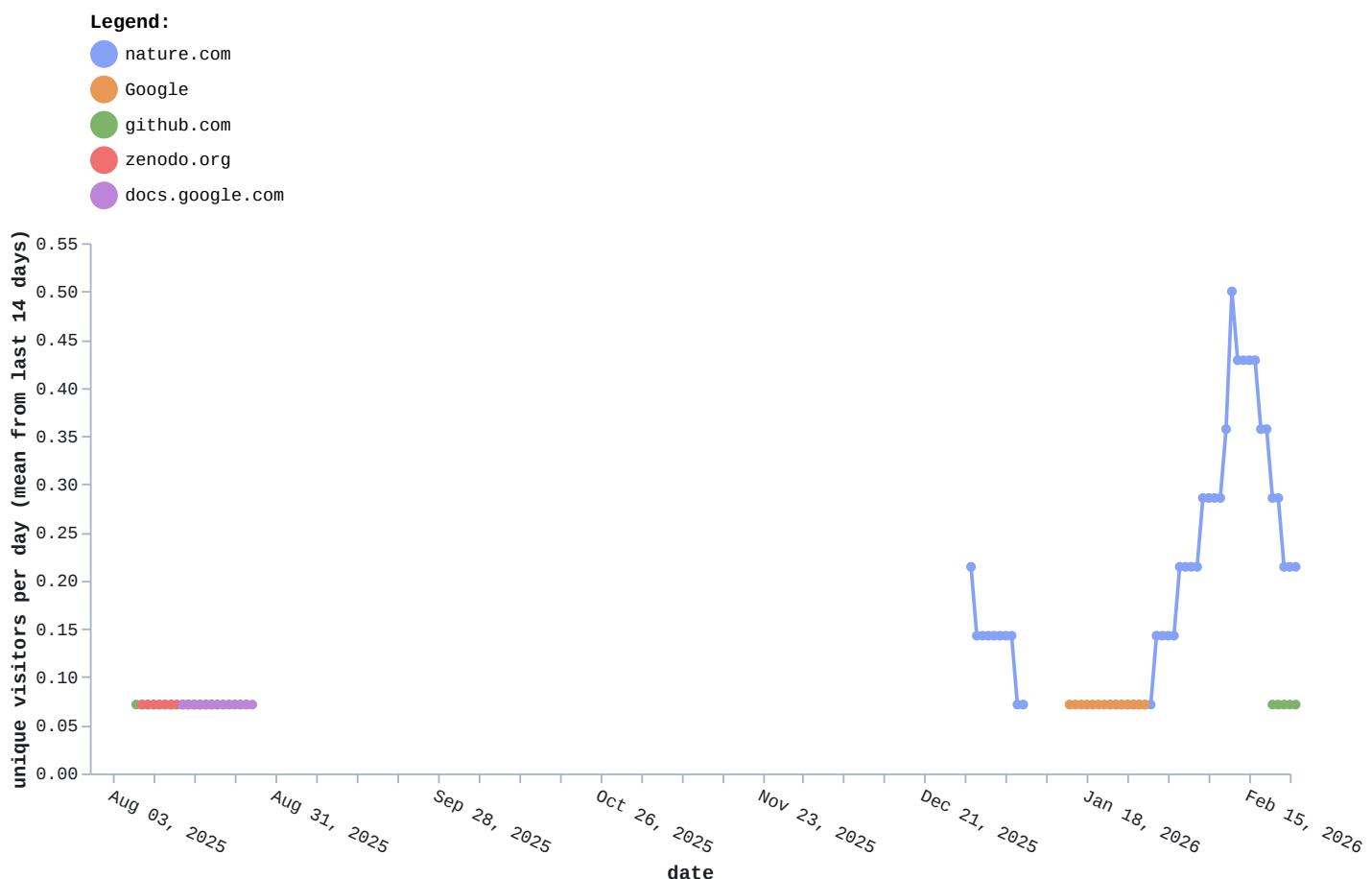
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This repository has no forks yet.

# 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

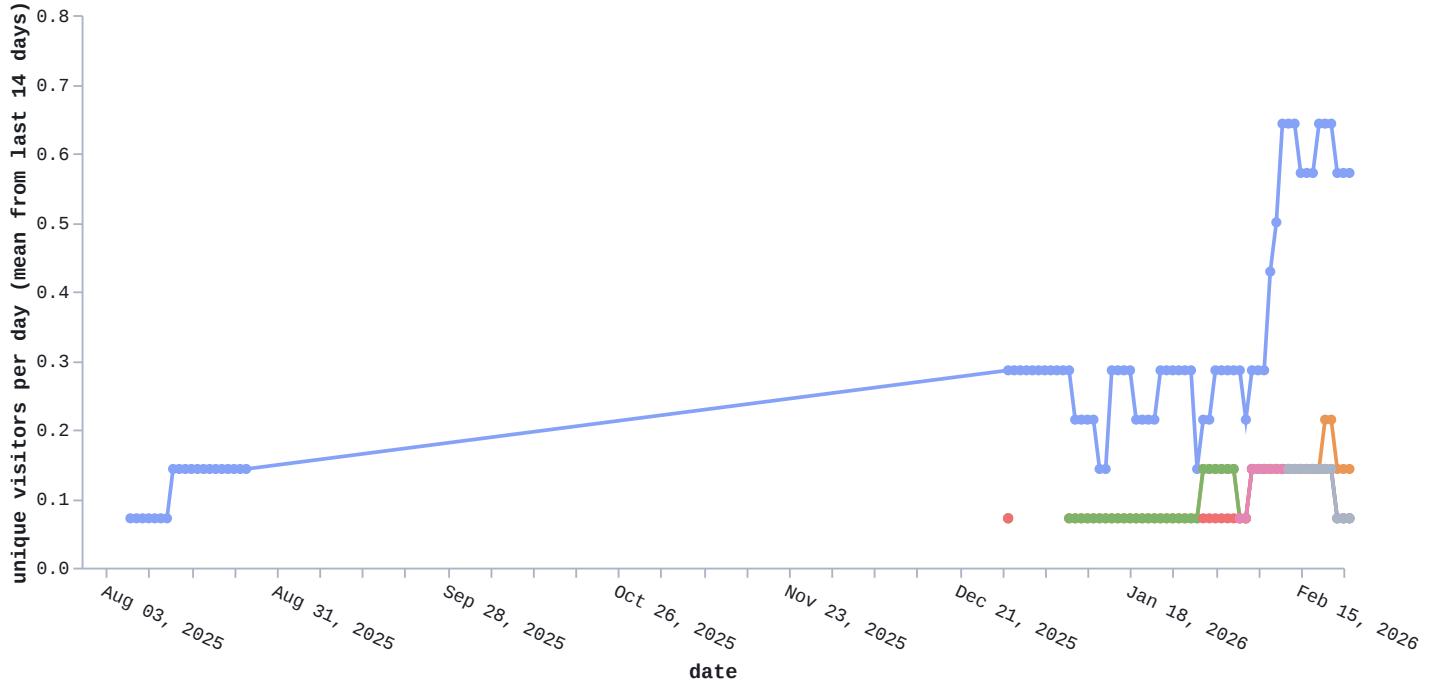


Top 15 referrers: 01: nature.com , 02: Google , 03: github.com , 04: zenodo.org , 05: docs.google.com

## Top paths

**Legend:**

- /
- /tree/main/01\_GWAS
- /blob/main/01\_GWAS/01\_UKB...
- /tree/main/01\_GWAS/02\_GWA...
- /blob/main/07\_SBayesS/03\_...
- /tree/main/06\_LDSC
- /blob/main/07\_SBayesS/03\_...



Top 15 paths: 01: / , 02: /tree/main/01\_GWAS , 03:

/blob/main/01\_GWAS/01\_UKB\_phenotype/UKB\_sex\_depression\_fastGWA\_final.html , 04:

/tree/main/01\_GWAS/02\_GWAS\_GxS\_example\_scripts , 05:

/blob/main/07\_SBayesS/03\_c\_SBayesS\_h2\_Blokland21\_male.sh , 06: /tree/main/06\_LDSC , 07:

/blob/main/07\_SBayesS/03\_a\_SBayesS\_h2.sh , 08: /tree/main/01\_GWAS/01\_UKB\_phenotype , 09:

/tree/main , 10: /tree/main/13\_Publication\_figures , 11: /tree/main/07\_SBayesS , 12:

/blob/main/08\_GWASpw/MDD\_BMI/femaleMDD\_femaleBMI/03\_genome\_wide\_cor.R , 13:

/blob/main/08\_GWASpw/femaleMDD\_maleMDD/05\_GWAS\_pw\_evidence\_models.sh , 14:

/blob/main/08\_GWASpw/femaleMDD\_maleMDD/06\_Plot\_Miami\_gwaspw.sh , 15:

/blob/main/01\_GWAS/02\_GWAS\_GxS\_example\_scripts/02\_a\_GCTA\_GRM\_autosomes.sh