

Resource Summary Report

Generated by T1D on Jul 31, 2026

TimeZone

RRID:SCR_018564

Type: Tool

Proper Citation

TimeZone (RRID:SCR_018564)

Resource Information

URL: <https://sourceforge.net/projects/timezone1/>

Proper Citation: TimeZone (RRID:SCR_018564)

Description: Software package to detect footprints of positive selection for functionally adaptive point mutations in microbial genomes.

Resource Type: data processing software, software application, data analysis software, software resource

Defining Citation: [PMID:23471110](https://pubmed.ncbi.nlm.nih.gov/23471110/)

Keywords: Detect footprint, positive selection, functionally adaptive, point mutation, microbial genome

Funding: NIGMS R01 GM084318;
NIAID RC4 AI092828

Availability: Free, Available for download, Freely available

Resource Name: TimeZone

Resource ID: SCR_018564

Record Creation Time: 20220129T080340+0000

Record Last Update: 20260731T043037+0000

Ratings and Alerts

No rating or validation information has been found for TimeZone.

No alerts have been found for TimeZone.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at [T1D](#) .

Robinson SM, et al. (2022) Distinct Potentially Adaptive Accumulation of Truncation Mutations in *Salmonella enterica* serovar Typhi and *Salmonella enterica* serovar Paratyphi A. *Microbiology spectrum*, 10(3), e0196921.