

Resource Summary Report

Generated by SPARC SAWG on Sep 18, 2026

TSSer

RRID:SCR_006419

Type: Tool

Proper Citation

TSSer (RRID:SCR_006419)

Resource Information

URL: <http://www.clipz.unibas.ch/downloads/TSSer/index.php>

Proper Citation: TSSer (RRID:SCR_006419)

Description: A computational pipeline to analyze differential RNA sequencing (dRNA-seq) data to determine transcription start sites genome-wide.

Abbreviations: TSSer

Synonyms: TSSer: a computational pipeline to identify transcription start sites in bacterial genomes

Resource Type: software resource

Defining Citation: [PMID:24371151](#)

Keywords: differential rna sequencing, transcription start site, rna-seq, genome, bio.tools

Availability: GNU General Public License

Resource Name: TSSer

Resource ID: SCR_006419

Alternate IDs: biotools:tsser, OMICS_02191

Alternate URLs: <https://bio.tools/tsser>

Record Creation Time: 20220129T080236+0000

Record Last Update: 20260912T195639+0000

Ratings and Alerts

No rating or validation information has been found for TSSer.

No alerts have been found for TSSer.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.