# Resource Summary Report

Generated by FDI Lab - SciCrunch.org on Mar 31, 2025

## ScriptManager

RRID:SCR_021797
Type: Tool

### Proper Citation

ScriptManager (RRID:SCR_021797)

### Resource Information

**URL:** https://github.com/CEGRcode/scriptmanager

**Proper Citation:** ScriptManager (RRID:SCR_021797)

**Description:** Interactive GUI Java software package for analysis of high throughput genomic sequencing data.

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

**Keywords:** High throughput genomic sequencing data, genomic sequencing data, NGS analysis scripts

**Funding:**

**Availability:** Free, Available for download, Freely available

**Resource Name:** ScriptManager

**Resource ID:** SCR_021797

**License:** MIT License

**Record Creation Time:** 20220129T080357+0000

**Record Last Update:** 20250331T061746+0000

### Ratings and Alerts

No rating or validation information has been found for ScriptManager.

No alerts have been found for ScriptManager.

## Data and Source Information

**Source:** SciCrunch Registry

## Usage and Citation Metrics

We found 6 mentions in open access literature.

**Listed below are recent publications.** The full list is available at FDI Lab - SciCrunch.org.

Louder RK, et al. (2024) Molecular basis of global promoter sensing and nucleosome capture by the SWR1 chromatin remodeler. Cell, 187(24), 6849.

van Breugel ME, et al. (2023) Locus-specific proteome decoding reveals Fpt1 as a chromatin-associated negative regulator of RNA polymerase III assembly. Molecular cell, 83(23), 4205.

John J, et al. (2022) Genome-wide promoter assembly in E. coli measured at single-base resolution. Genome research, 32(5), 878.

Mittal C, et al. (2022) An integrated SAGA and TFIID PIC assembly pathway selective for poised and induced promoters. Genes & development, 36(17-18), 985.

Zhao T, et al. (2021) Ssl2/TFIIH function in transcription start site scanning by RNA polymerase II in Saccharomyces cerevisiae. eLife, 10.

Lai WKM, et al. (2021) A ChIP-exo screen of 887 Protein Capture Reagents Program transcription factor antibodies in human cells. Genome research, 31(9), 1663.