

# Resource Summary Report

Generated by [PRECISE-TBI](#) on Aug 1, 2026

## Squeezambler

RRID:SCR\_004385

Type: Tool

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### Proper Citation

Squeezambler (RRID:SCR\_004385)

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### Resource Information

**URL:** <http://compbio.cs.wayne.edu/software/squeezambler/>

**Proper Citation:** Squeezambler (RRID:SCR\_004385)

**Description:** Software to sequence and de novo assemble all distinct genomes present in a microbial sample with a sequencing cost and computational complexity proportional to the number of genome types, rather than the number of cells.

**Abbreviations:** Squeezambler

**Resource Type:** software resource

**Defining Citation:** [PMID:23918251](#)

**Resource Name:** Squeezambler

**Resource ID:** SCR\_004385

**Alternate IDs:** OMICS\_01503

**Record Creation Time:** 20220129T080224+0000

**Record Last Update:** 20260725T190552+0000

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### Ratings and Alerts

No rating or validation information has been found for Squeezambler.

No alerts have been found for Squeezambler.

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### Data and Source Information

**Source:** [SciCrunch Registry](#)

## Usage and Citation Metrics

We have not found any literature mentions for this resource.