**Squeezambler**

**RRID:** SCR_004385  
**Type:** Tool

**Proper Citation**

Squeezambler (RRID:SCR_004385)

**Resource Information**

**URL:** [http://compbio.cs.wayne.edu/software/squeezambler/](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](https://www.ncbi.nlm.nih.gov/pubmed/23918251)

**Resource Name:** Squeezambler

**Resource ID:** SCR_004385

**Alternate IDs:** OMICS_01503

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

**Record Last Update:** 20240424T182803+0000

**Ratings and Alerts**

No rating or validation information has been found for Squeezambler.

No alerts have been found for Squeezambler.
Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We have not found any literature mentions for this resource.