# **Resource Summary Report**

Generated by NIF on Apr 17, 2025

# **mubiomics**

RRID:SCR\_006785

Type: Tool

## **Proper Citation**

mubiomics (RRID:SCR\_006785)

#### **Resource Information**

URL: http://sourceforge.net/projects/mubiomics/

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**Description:** A set of scripts (mostly python) for processing reads generated by the Roche 454 or Illumina next-gen sequencing platforms. Included are quality control, read demultiplexing and microbiome characterisation scripts for use with usearch, pplacer and RDP classifier.

Abbreviations: mubiomics

Synonyms: mubiomics - Scripts for processing next-gen sequencing data

**Resource Type:** software resource

**Funding:** 

**Resource Name:** mubiomics

Resource ID: SCR 006785

Alternate IDs: OMICS\_01059

Record Creation Time: 20220129T080238+0000

Record Last Update: 20250410T065453+0000

### **Ratings and Alerts**

No rating or validation information has been found for mubiomics.

No alerts have been found for mubiomics.

### **Data and Source Information**

Source: SciCrunch Registry

## **Usage and Citation Metrics**

We found 3 mentions in open access literature.

**Listed below are recent publications.** The full list is available at NIF.

Li Y, et al. (2022) Amlodipine, an anti-hypertensive drug, alleviates non-alcoholic fatty liver disease by modulating gut microbiota. British journal of pharmacology, 179(9), 2054.

van Eeden C, et al. (2018) V2-Directed Vaccine-like Antibodies from HIV-1 Infection Identify an Additional K169-Binding Light Chain Motif with Broad ADCC Activity. Cell reports, 25(11), 3123.

Smith BC, et al. (2012) The cervical microbiome over 7 years and a comparison of methodologies for its characterization. PloS one, 7(7), e40425.