# **Resource Summary Report**

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### rtax

RRID:SCR\_024320

Type: Tool

### **Proper Citation**

rtax (RRID:SCR\_024320)

#### **Resource Information**

URL: https://github.com/davidsoergel/rtax/

**Proper Citation:** rtax (RRID:SCR\_024320)

**Description:** Software tool for rapid and accurate taxonomic classification of short pairedend sequence reads from the 16S ribosomal RNA gene.

Resource Type: software resource, software application

**Defining Citation: PMID:22237546** 

**Keywords:** taxonomic classification, short paired-end sequence reads, 16S ribosomal RNA gene,

**Funding:** 

Availability: Free, Available for download, Freely available,

Resource Name: rtax

Resource ID: SCR\_024320

Alternate IDs: OMICS\_28979

Alternate URLs: https://sources.debian.org/src/rtax/

**Record Creation Time:** 20230830T050217+0000

Record Last Update: 20250421T054527+0000

# Ratings and Alerts

No rating or validation information has been found for rtax.

No alerts have been found for rtax.

### Data and Source Information

Source: SciCrunch Registry

# **Usage and Citation Metrics**

We have not found any literature mentions for this resource.