

Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 8, 2026

TACOA

RRID:SCR_000107

Type: Tool

Proper Citation

TACOA (RRID:SCR_000107)

Resource Information

URL: <http://www.cebitec.uni-bielefeld.de/index.php/2-uncategorised/99-tacoa?highlight=WyJOYWNVYSJd>

Proper Citation: TACOA (RRID:SCR_000107)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on July 31,2025. Software that can accurately predict the taxonomic origin of genomic fragments from metagenomic data sets by combining the advantages of the k -NN approach with a smoothing kernel function.

Abbreviations: TACOA

Resource Type: software resource

Defining Citation: [PMID:19210774](#)

Keywords: taxonomy, genomic fragment, metagenome

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: TACOA

Resource ID: SCR_000107

Alternate IDs: OMICS_01467

Record Creation Time: 20220129T080159+0000

Record Last Update: 20260905T132413+0000

Ratings and Alerts

No rating or validation information has been found for TACOA.

No alerts have been found for TACOA.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.