

# Resource Summary Report

Generated by T1D on Aug 2, 2026

## Bio-Tools-Run-Alignment-TCoffee

RRID:SCR\_024070

Type: Tool

---

### Proper Citation

Bio-Tools-Run-Alignment-TCoffee (RRID:SCR\_024070)

---

### Resource Information

**URL:** <https://metacpan.org/dist/Bio-Tools-Run-Alignment-TCoffee>

**Proper Citation:** Bio-Tools-Run-Alignment-TCoffee (RRID:SCR\_024070)

**Description:** Software object for calculation of multiple sequence alignment from set of unaligned sequences or alignments using the TCoffee program.

**Synonyms:** libbio-tools-run-alignment-tcoffee-perl

**Resource Type:** software resource, software application

**Keywords:** calculation of multiple sequence alignment, unaligned sequences set, alignments set, TCoffee program,

**Availability:** Free, Available for download, Freely available,

**Resource Name:** Bio-Tools-Run-Alignment-TCoffee

**Resource ID:** SCR\_024070

**Alternate URLs:** <https://sources.debian.org/src/libbio-tools-run-alignment-tcoffee-perl/>

**License:** perl\_5

**Record Creation Time:** 20230824T050211+0000

**Record Last Update:** 20260801T191123+0000

---

### Ratings and Alerts

No rating or validation information has been found for Bio-Tools-Run-Alignment-TCoffee.

No alerts have been found for Bio-Tools-Run-Alignment-TCoffee.

---

## Data and Source Information

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

---

## Usage and Citation Metrics

We have not found any literature mentions for this resource.