**T-Coffee**

**RRID:** SCR_011818  
**Type:** Tool

**Proper Citation**

T-Coffee (RRID:SCR_011818)

---

**Resource Information**


**Proper Citation:** T-Coffee (RRID:SCR_011818)

**Description:** A multiple sequence alignment server which can align Protein, DNA and RNA sequences.

**Resource Type:** Resource, analysis service resource, data analysis service, service resource, production service resource

**References:** PMID:10964570

**Keywords:** bio.tools

**Parent Organization:** CRG; Barcelona; Spain

**Website Status:** Last checked up

**Abbreviations:** T-Coffee

**Resource Name:** T-Coffee

**Resource ID:** SCR_011818

**Alternate IDs:** OMICS_00989, biotools:tcoffee

**Alternate URLs:** [https://bio.tools/tcoffee](https://bio.tools/tcoffee)

---

Ratings and Alerts
No rating or validation information has been found for T-Coffee.

No alerts have been found for T-Coffee.

Data and Source Information

**Source:** SciCrunch Registry

Usage and Citation Metrics

We found 644 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [FDI Lab - SciCrunch.org](https://www.FDI-Lab.org).


- **Wood D, et al. (2020)** Phylogenomics indicates the "living fossil" Isoetes diversified in the Cenozoic. PloS one, 15(6), e0227525.

- **Signetti L, et al. (2020)** Inhibition of Patched Drug Efflux Increases Vemurafenib Effectiveness against Resistant BrafMelanoma. Cancers, 12(6).


- **Jensen MM, et al. (2020)** The origins and developments of sulfation-prone tyrosine-rich and acidic N- and C-terminal extensions of class II and III small leucine-rich repeat proteins shed light on connective tissue evolution in vertebrates. BMC evolutionary biology, 20(1), 73.


- **Gazi MA, et al. (2020)** Questing functions and structures of hypothetical proteins from


