

Resource Summary Report

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TeamTat

RRID:SCR_023439

Type: Tool

Proper Citation

TeamTat (RRID:SCR_023439)

Resource Information

URL: <https://www.teamtat.org>

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Description: Web based collaborative text annotation tool. Used for managing multi-user, multi-label document annotation. Project managers can specify annotation schema for entities and relations and select annotators and distribute documents anonymously to prevent bias. Document input format can be plain text, PDF or BioC (uploaded locally or automatically retrieved from PubMed/PMC), and output format is BioC with inline annotations. Displays figures from full text.

Resource Type: web application, software resource

Defining Citation: [PMID:32383756](#)

Keywords: Text annotation, team collaboration, collaborative text annotation,

Funding: MSIT ;
NIH

Availability: Free, Freely available

Resource Name: TeamTat

Resource ID: SCR_023439

Alternate URLs: <https://github.com/ncbi-nlp/TeamTat>

License: MIT License

Record Creation Time: 20230405T050216+0000

Record Last Update: 20260728T043657+0000

Ratings and Alerts

No rating or validation information has been found for TeamTat.

No alerts have been found for TeamTat.

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 [dkNET](#) .

Lai PT, et al. (2024) EnzChemRED, a rich enzyme chemistry relation extraction dataset. Scientific data, 11(1), 982.

Vollmar M, et al. (2024) Dataset from a human-in-the-loop approach to identify functionally important protein residues from literature. Scientific data, 11(1), 1032.

Neves M, et al. (2023) Automatic classification of experimental models in biomedical literature to support searching for alternative methods to animal experiments. Journal of biomedical semantics, 14(1), 13.