

Resource Summary Report

Generated by RRID on Aug 6, 2026

Django

RRID:SCR_012855

Type: Tool

Proper Citation

Django (RRID:SCR_012855)

Resource Information

URL: <http://www.djangoproject.com/>

Proper Citation: Django (RRID:SCR_012855)

Description: Django is a high-level Python Web framework that encourages rapid development and clean, pragmatic design. Developed four years ago by a fast-moving online-news operation, Django was designed to handle two challenges: the intensive deadlines of a newsroom and the stringent requirements of the experienced Web developers who wrote it. It lets you build high-performing, elegant Web applications quickly. Django focuses on automating as much as possible and adhering to the DRY principle.

Synonyms: Django

Resource Type: software resource

Resource Name: Django

Resource ID: SCR_012855

Alternate IDs: nif-0000-30460

Record Creation Time: 20220129T080312+0000

Record Last Update: 20260801T190452+0000

Ratings and Alerts

No rating or validation information has been found for Django.

No alerts have been found for Django.

Data and Source Information

Usage and Citation Metrics

We found 561 mentions in open access literature.

Listed below are recent publications. The full list is available at [RRID](#) .

Hadarovich A, et al. (2026) PICNIC web server for predicting proteins involved in biomolecular condensates. *Bioinformatics* (Oxford, England), 42(1).

Barca L, et al. (2025) Emotional distress and affective knowledge representation one year after the COVID-19 outbreak. *PloS one*, 20(1), e0311009.

Miao Z, et al. (2025) Interactive visualization and interpretation of pangenome graphs by linear reference-based coordinate projection and annotation integration. *Genome research*, 35(2), 296.

Feng Q, et al. (2025) A membrane permeability database for nonpeptidic macrocycles. *Scientific data*, 12(1), 10.

Liu J, et al. (2025) Replacing manual planning with automatic iterative planning for locally advanced rectal cancer VMAT treatment. *Journal of applied clinical medical physics*, 26(1), e14552.

Lan J, et al. (2025) MoLPre: A Machine Learning Model to Predict Metastasis of cT1 Solid Lung Cancer. *Clinical and translational science*, 18(4), e70186.

Stanley DN, et al. (2025) ADAPT-marketplace - A web-application for XML-based template sharing in radiation oncology. *Journal of applied clinical medical physics*, 26(7), e70148.

Wang J, et al. (2025) hAb-Convergent: an antibody rearrangement analysis system for therapeutic antibody engineering based on convergent evolution. *Nucleic acids research*, 53(W1), W297.

Mo O, et al. (2025) mRNAdesigner: an integrated web server for optimizing mRNA design and protein translation in eukaryotes. *Nucleic acids research*, 53(W1), W415.

Kumar P, et al. (2025) MechBERT: Language Models for Extracting Chemical and Property Relationships about Mechanical Stress and Strain. *Journal of chemical information and modeling*, 65(4), 1873.

Silawarawet K, et al. (2025) Heat Stroke Warning System Prototype for Athletes: A Pilot Study. *Sensors* (Basel, Switzerland), 25(2).

Bai H, et al. (2025) TetraRNA, a tetra-class machine learning model for deciphering the coding potential derivation of RNA world. *Computational and structural biotechnology journal*, 27, 1305.

Rigutto G, et al. (2025) Mapping assays to the key characteristics of carcinogens to support decision-making. Database : the journal of biological databases and curation, 2025.

Joshi J, et al. (2025) DICED (Database of Identified Cleavage Sites Endemic to Diseases States): A Searchable Web Interface for Terminomics/Degradomics. Proteomics, 25(13), e202500007.

Ralf A, et al. (2025) UYSD: a novel data repository accessible via public website for worldwide population frequencies of Y-SNP haplogroups. European journal of human genetics : EJHG, 33(7), 904.

Mareuil F, et al. (2025) InDeepNet: a web platform for predicting functional binding sites in proteins using InDeep. Nucleic acids research, 53(W1), W324.

Battistel C, et al. (2025) MoPSeq-DB: a user-friendly web application for genomic data management and analysis of marine mollusc pathogens. Database : the journal of biological databases and curation, 2025.

??kuta C, et al. (2025) ECBD: European chemical biology database. Nucleic acids research, 53(D1), D1383.

Meng X, et al. (2025) GTO: a comprehensive gene therapy omnibus. Nucleic acids research, 53(D1), D1393.

Liu Z, et al. (2025) SolR: a comprehensive Solanaceae information resource for comparative and functional genomic study. Nucleic acids research, 53(D1), D1623.