

Resource Summary Report

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Gutentag

RRID:SCR_003051

Type: Tool

Proper Citation

Gutentag (RRID:SCR_003051)

Resource Information

URL: <https://code.google.com/p/gutentag/>

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Description: An interactive, user-editable genetic sequence database tool, targeted at molecular biology research groups that can be browsed using tags. The tool is Web 2.0-flavoured, allowing users to do more than just retrieve information. Its focus on user-editability is supported by the use of tags (metadata) associated with genetic sequences. Several methods of retrieving stored data are available including tag-clouds, BLAST and keyword searches. Also, sequence tags related to HGNC gene names, conserved domains (CDD) and GO terms can be automatically generated given sequence data. The tool is constructed using the high-level Python web framework, Django, with a SQLite3 backend.

Synonyms: gutentag - A genetic sequence database for labs that can be browsed using tags

Resource Type: software resource

Defining Citation: [PMID:14640709](#)

Keywords: standalone software, python, django, genetics, sequence

Funding:

Availability: GNU General Public License, v3

Resource Name: Gutentag

Resource ID: SCR_003051

Alternate IDs: OMICS_02396

Record Creation Time: 20220129T080216+0000

Record Last Update: 20250420T014134+0000

Ratings and Alerts

No rating or validation information has been found for Gutentag.

No alerts have been found for Gutentag.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.