

# Resource Summary Report

Generated by [dkNET](#) on Jul 30, 2026

## Gutentag

RRID:SCR\_003051

Type: Tool

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### Proper Citation

Gutentag (RRID:SCR\_003051)

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### Resource Information

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

**Proper Citation:** Gutentag (RRID:SCR\_003051)

**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

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

**Resource Name:** Gutentag

**Resource ID:** SCR\_003051

**Alternate IDs:** OMICS\_02396

**Record Creation Time:** 20220129T080216+0000

**Record Last Update:** 20260725T190537+0000

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## Ratings and Alerts

No rating or validation information has been found for Gutentag.

No alerts have been found for Gutentag.

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## Data and Source Information

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

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.