

Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 8, 2026

FIGG

RRID:SCR_012064

Type: Tool

Proper Citation

FIGG (RRID:SCR_012064)

Resource Information

URL: <http://insilicogenome.sourceforge.net/>

Proper Citation: FIGG (RRID:SCR_012064)

Description: A large-scale whole genome simulation tool which generates large numbers of whole genomes with known sequence characteristics based on direct sampling of experimentally known or theorized variations.

Synonyms: Frequency-based Insilico Genome Generator

Resource Type: software resource

Defining Citation: [PMID:24885193](#)

Keywords: standalone software, unix/linux, mac os x, windows, java, mapreduce

Availability: Free, Public

Resource Name: FIGG

Resource ID: SCR_012064

Alternate IDs: OMICS_04164

Record Creation Time: 20220129T080308+0000

Record Last Update: 20260905T132719+0000

Ratings and Alerts

No rating or validation information has been found for FIGG.

No alerts have been found for FIGG.

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 [PRECISE-TBI](#) .

Guerrini CJ, et al. (2025) FIGG at 5: An update on U.S. public perspectives on forensic investigative genetic genealogy five years after its introduction to criminal investigations. Forensic science international, 367, 112372.

Budowle B, et al. (2023) A cost-benefit analysis for use of large SNP panels and high throughput typing for forensic investigative genetic genealogy. International journal of legal medicine, 137(5), 1595.

Woudenberg T, et al. (2019) Additional Evidence on Serological Correlates of Protection against Measles: An Observational Cohort Study among Once Vaccinated Children Exposed to Measles. Vaccines, 7(4).