**DWGSIM**

**RRID:** SCR_002342  
**Type:** Tool

**Proper Citation**

DWGSIM (RRID:SCR_002342)

**Resource Information**

**URL:** [https://github.com/nh13/DWGSIM](https://github.com/nh13/DWGSIM)  
**Proper Citation:** DWGSIM (RRID:SCR_002342)  
**Description:** Whole Genome Simulator for Next-Generation Sequencing.  
**Abbreviations:** DWGSIM  
**Resource Type:** software resource  
**Keywords:** next-generation sequencing, whole genome simulation  
**Availability:** GNU General Public License, v2  
**Resource Name:** DWGSIM  
**Resource ID:** SCR_002342  
**Alternate IDs:** OMICS_00249  
**Alternate URLs:** [https://sources.debian.org/src/dwgsim/](https://sources.debian.org/src/dwgsim/)  
**Record Creation Time:** 20220129T080212+0000  
**Record Last Update:** 20240424T182739+0000

**Ratings and Alerts**

No rating or validation information has been found for DWGSIM.
No alerts have been found for DWGSIM.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 49 mentions in open access literature.

Listed below are recent publications. The full list is available at FDI Lab - SciCrunch.org.


Chen J, et al. (2024) CropGS-Hub: a comprehensive database of genotype and phenotype resources for genomic prediction in major crops. Nucleic acids research, 52(D1), D1519.

Wang S, et al. (2023) SpecHLA enables full-resolution HLA typing from sequencing data. Cell reports methods, 3(9), 100589.


