

Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 2026

forqs

RRID:SCR_000643

Type: Tool

Proper Citation

forqs (RRID:SCR_000643)

Resource Information

URL: <https://bitbucket.org/dkessner/forqs>

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Description: Software for forward-in-time population genetics simulation that tracks individual haplotype chunks as they recombine each generation. It also also models quantitative traits and selection on those traits.

Abbreviations: forqs

Synonyms: Forward-in-time simulation of Recombination, and Selection, Quantitative traits

Resource Type: software application, simulation software, software resource

Defining Citation: [PMID:24336146](#)

Keywords: c++, linux, osx, windows, command line, simulation, recombination, quantitative trait, selection, haplotype pattern

Funding: NHGRI HG002536;
NHGRI R01 HG007089;
NSF EF-0928690

Availability: Free, Available for download, Freely available

Resource Name: forqs

Resource ID: SCR_000643

Alternate IDs: OMICS_02196

License: BSD License

Record Creation Time: 20220129T080202+0000

Record Last Update: 20260728T043100+0000

Ratings and Alerts

No rating or validation information has been found for forqs.

No alerts have been found for forqs.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.