

Resource Summary Report

Generated by [nidm-terms](#) on Jul 30, 2026

OLORIN

RRID:SCR_002015

Type: Tool

Proper Citation

OLORIN (RRID:SCR_002015)

Resource Information

URL: <http://www.sanger.ac.uk/science/tools/olorin>

Proper Citation: OLORIN (RRID:SCR_002015)

Description: An interactive filtering tool for next generation sequencing data coming from the study of large complex disease pedigrees. It integrates gene flow output from Merlin and next generation sequencing data. Users can interactively filter and prioritize variants based on haplotype sharing across different sets of selected individuals and allele frequency in reference datasets. (entry from Genetic Analysis Software)

Abbreviations: OLORIN

Resource Type: software application, software resource

Defining Citation: [PMID:23052039](#)

Keywords: gene, genetic, genomic, java, any platform with java 1.6 or later, next generation sequencing, variant, haplotype, allele frequency, java swing

Availability: Free, Available for download, Freely available

Resource Name: OLORIN

Resource ID: SCR_002015

Alternate IDs: nlx_154503, OMICS_01556

Alternate URLs: <http://sourceforge.net/p/olorin/>

License: MIT License

Record Creation Time: 20220129T080211+0000

Record Last Update: 20260727T040312+0000

Ratings and Alerts

No rating or validation information has been found for OLORIN.

No alerts have been found for OLORIN.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.