

Resource Summary Report

Generated by SPARC SAWG on Aug 10, 2026

RADtools

RRID:SCR_000324

Type: Tool

Proper Citation

RADtools (RRID:SCR_000324)

Resource Information

URL: <https://github.com/johnomics/RADtools>

Proper Citation: RADtools (RRID:SCR_000324)

Description: Software tools for processing RAD Sequencing Illumina reads. Version 1.0 is a pipeline for transforming Illumina reads into candidate genetic markers.

Resource Type: software resource

Defining Citation: [PMID:21541297](https://pubmed.ncbi.nlm.nih.gov/21541297/)

Keywords: standalone software, illumina

Availability: Free, Available for download, Freely available

Resource Name: RADtools

Resource ID: SCR_000324

Alternate IDs: OMICS_03727

Alternate URLs: <https://www.wiki.ed.ac.uk/display/RADSequencing/>

License: GNU General Public License, v3

Record Creation Time: 20220129T080200+0000

Record Last Update: 20260808T185719+0000

Ratings and Alerts

No rating or validation information has been found for RADtools.

No alerts have been found for RADtools.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at [SPARC SAWG](#) .

Gamble T, et al. (2017) The Discovery of XY Sex Chromosomes in a Boa and Python.
Current biology : CB, 27(14), 2148.