Resource Summary Report

Generated by <u>RRID</u> on Apr 28, 2025

parSMURF

RRID:SCR_017560 Type: Tool

Proper Citation

parSMURF (RRID:SCR_017560)

Resource Information

URL: https://github.com/AnacletoLAB/parSMURF

Proper Citation: parSMURF (RRID:SCR_017560)

Description: Open source software package as high performance computing imbalance aware machine learning tool for genome wide detection of pathogenic variants.

Resource Type: data processing software, software application, software resource

Keywords: High, performance, computing, imbalance, aware, machine, learning, genome, wide, detection, pathogenic, variant, bio.tools

Funding:

Availability: Free, Available for download, Freely available

Resource Name: parSMURF

Resource ID: SCR_017560

Alternate IDs: biotools:parsmurf

Alternate URLs: https://bio.tools/parsmurf

License: GNU General Public License v3.0

Record Creation Time: 20220129T080335+0000

Record Last Update: 20250428T054055+0000

Ratings and Alerts

No rating or validation information has been found for parSMURF.

No alerts have been found for parSMURF.

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 <u>RRID</u>.

Petrini A, et al. (2020) parSMURF, a high-performance computing tool for the genome-wide detection of pathogenic variants. GigaScience, 9(5).