

Resource Summary Report

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Benchmark of human deep intronic variation

RRID:SCR_023315

Type: Tool

Proper Citation

Benchmark of human deep intronic variation (RRID:SCR_023315)

Resource Information

URL: https://github.com/PedroBarbosa/DeepIntronic_Benchmark

Proper Citation: Benchmark of human deep intronic variation (RRID:SCR_023315)

Description: Repository of code, data, and documentation to reproduce computational evaluation performed to identify the best tools/models to predict genetic variation in deep intronic regions of human genome.

Synonyms: Benchmark of deep intronic variation

Resource Type: software resource

Defining Citation: [DOI:10.1101/2023.02.17.528928](https://doi.org/10.1101/2023.02.17.528928)

Keywords: reproduce computational evaluation, predict genetic variation, deep intronic regions, human genome

Availability: Free, Available for download, Freely available

Resource Name: Benchmark of human deep intronic variation

Resource ID: SCR_023315

License: GNU GPL v3.0

Record Creation Time: 20230303T050211+0000

Record Last Update: 20260808T190244+0000

Ratings and Alerts

No rating or validation information has been found for Benchmark of human deep intronic variation.

No alerts have been found for Benchmark of human deep intronic variation.

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](#) .

Fortier N, et al. (2026) Analyzing the performance of deep learning splice prediction algorithms. PloS one, 21(5), e0348885.