

Resource Summary Report

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Prepare_SplicingPredictors

RRID:SCR_023316

Type: Tool

Proper Citation

Prepare_SplicingPredictors (RRID:SCR_023316)

Resource Information

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

Proper Citation: Prepare_SplicingPredictors (RRID:SCR_023316)

Description: Software tool as set of scripts to prepare the input for several sequence-based RNA splicing methods from VCF files.

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

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

Keywords: RNA splicing, sequence based models, prepare input from VCF files,

Funding: Fundação para a Ciência e a Tecnologia (FCT) Portugal

Availability: Free, Available for download, Freely available

Resource Name: Prepare_SplicingPredictors

Resource ID: SCR_023316

License: GNU GPL v3.0

Record Creation Time: 20230303T050211+0000

Record Last Update: 20260813T055301+0000

Ratings and Alerts

No rating or validation information has been found for Prepare_SplicingPredictors.

No alerts have been found for Prepare_SplicingPredictors.

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 [RRID](#) .

Barbosa P, et al. (2022) Computational prediction of human deep intronic variation. GigaScience, 12.