

Resource Summary Report

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TranscriptAchilles

RRID:SCR_016849

Type: Tool

Proper Citation

TranscriptAchilles (RRID:SCR_016849)

Resource Information

URL: <http://biotecnun.unav.es:8080/app/TranscriptAchilles>

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Description: Software genome-wide tool to predict transcript biomarkers of gene essentiality in cancer. This tool can be used to predict new potential target genes with their corresponding biomarkers (either transcript or gene expression).

Abbreviations: TranscriptAchilles

Resource Type: sequence analysis software, software resource, data analysis software, data processing software, web application, software application

Keywords: predict, transcript, biomarker, gene, inhibition, isoform, cancer, cell line, expression, essentiality, target gene

Availability: Free, Available for download, Freely available

Resource Name: TranscriptAchilles

Resource ID: SCR_016849

Alternate URLs: <https://gitlab.com/fcarazo.m/transcriptachilles/>

License: GPLv3

Record Creation Time: 20220129T080332+0000

Record Last Update: 20260807T042858+0000

Ratings and Alerts

No rating or validation information has been found for TranscriptAchilles.

No alerts have been found for TranscriptAchilles.

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](#) .

Carazo F, et al. (2019) TranscriptAchilles: a genome-wide platform to predict isoform biomarkers of gene essentiality in cancer. GigaScience, 8(4).