Resource Summary Report

Generated by dkNET on Apr 30, 2025

FusionFinder

RRID:SCR_011894
Type: Tool

Proper Citation

FusionFinder (RRID:SCR_011894)

Resource Information

URL: http://bioinformatics.childhealthresearch.org.au/software/fusionfinder/

Proper Citation: FusionFinder (RRID:SCR_011894)

Description: A perl-based software package, which can be used to find fusion transcript candidates in RNA-Seq data.

Abbreviations: FusionFinder

Synonyms: FusionHunter: identifying fusion transcripts using paired-end RNA-seq

Resource Type: software resource

Funding:

Resource Name: FusionFinder

Resource ID: SCR_011894

Alternate IDs: OMICS_01349

Record Creation Time: 20220129T080307+0000

Record Last Update: 20250420T014601+0000

Ratings and Alerts

No rating or validation information has been found for FusionFinder.

No alerts have been found for FusionFinder.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 4 mentions in open access literature.

Listed below are recent publications. The full list is available at dkNET.

Latysheva NS, et al. (2016) Discovering and understanding oncogenic gene fusions through data intensive computational approaches. Nucleic acids research, 44(10), 4487.

Panagopoulos I, et al. (2015) Novel KAT6B-KANSL1 fusion gene identified by RNA sequencing in retroperitoneal leiomyoma with t(10;17)(q22;q21). PloS one, 10(1), e0117010.

Panagopoulos I, et al. (2014) The "grep" command but not FusionMap, FusionFinder or ChimeraScan captures the CIC-DUX4 fusion gene from whole transcriptome sequencing data on a small round cell tumor with t(4;19)(q35;q13). PloS one, 9(6), e99439.

Francis RW, et al. (2012) FusionFinder: a software tool to identify expressed gene fusion candidates from RNA-Seq data. PloS one, 7(6), e39987.