Resource Summary Report

Generated by dkNET on May 20, 2025

nFuse

RRID:SCR_000066
Type: Tool

Proper Citation

nFuse (RRID:SCR_000066)

Resource Information

URL: https://code.google.com/p/nfuse/

Proper Citation: nFuse (RRID:SCR_000066)

Description: Software that predicts fusion transcripts and associated CGRs from matched RNA-seq and Whole Genome Shotgun Sequencing (WGSS).

Abbreviations: nFuse

Synonyms: nFuse: Discovery of Complex Genomic Rearrangements in Cancer

Resource Type: software resource

Defining Citation: PMID:22745232

Keywords: cancer, genomics

Related Condition: Cancer

Funding:

Availability: GNU General Public License, v3

Resource Name: nFuse

Resource ID: SCR_000066

Alternate IDs: OMICS_01353

Record Creation Time: 20220129T080159+0000

Record Last Update: 20250519T203042+0000

Ratings and Alerts

No rating or validation information has been found for nFuse.

No alerts have been found for nFuse.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Kumar S, et al. (2016) Comparative assessment of methods for the fusion transcripts detection from RNA-Seq data. Scientific reports, 6, 21597.

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