Resource Summary Report

Generated by RRID on Apr 19, 2025

NovelSeq

RRID:SCR_003136
Type: Tool

Proper Citation

NovelSeq (RRID:SCR_003136)

Resource Information

URL: http://compbio.cs.sfu.ca/software-novelseq

Proper Citation: NovelSeq (RRID:SCR_003136)

Description: Software pipeline to detect novel sequence insertions using high throughput paired-end whole genome sequencing data.

Abbreviations: NovelSeq

Synonyms: NovelSeq: Novel Sequence Insertion Detection

Resource Type: software resource

Defining Citation: PMID:20385726

Keywords: sequence, insertion, genome sequencing, genome, next-generation sequencing, illumina, unix, linux, c, bio.tools

Funding:

Resource Name: NovelSeq

Resource ID: SCR_003136

Alternate IDs: biotools:novelseq, nlx_156791, OMICS_02164

Alternate URLs: http://novelseq.sourceforge.net/, https://bio.tools/novelseq

Record Creation Time: 20220129T080217+0000

Record Last Update: 20250410T064943+0000

Ratings and Alerts

No rating or validation information has been found for NovelSeq.

No alerts have been found for NovelSeq.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We have not found any literature mentions for this resource.