Resource Summary Report

FreClu

RRID:SCR_005524
Type: Tool

Proper Citation

FreClu (RRID:SCR_005524)

Resource Information

URL: http://mlab.cb.k.u-tokyo.ac.jp/~quwei/DeNovoShortReadClustering/

Proper Citation: FreClu (RRID:SCR_005524)

Description: An algorithm for efficient frequency-based de novo short read clustering for error trimming in next-generation sequencing. It organizes erroneous short sequences originating in a single abundant sequence into a tree structure such that each child sequence is considered to be derived stochastically from its more abundant parent sequence because of sequencing errors.

Abbreviations: FreClu

Synonyms: FreClu - Efficient Frequency-based de Novo Short Read Clustering

Resource Type: software resource

Keywords: next-generation sequencing

Funding:

Resource Name: FreClu

Resource ID: SCR_005524

Alternate IDs: OMICS_01048

Record Creation Time: 20220129T080230+0000

Record Last Update: 20250410T065304+0000

Ratings and Alerts

No rating or validation information has been found for FreClu.

No alerts have been found for FreClu.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We have not found any literature mentions for this resource.