IDBA-UD
RRID:SCR_011912
Type: Tool

Proper Citation

IDBA-UD (RRID:SCR_011912)

Resource Information

**URL:** http://i.cs.hku.hk/~alse/hkubrg/projects/idba_ud/

**Proper Citation:** IDBA-UD (RRID:SCR_011912)

**Description:** Software for an iterative De Bruijn Graph De Novo Assembler for Short Reads Sequencing data with Highly Uneven Sequencing Depth.

**Resource Type:** Resource, software resource

**References:** PMID:22495754

**Parent Organization:** University of Hong Kong; Hong Kong; China

**Related resources:** Meta-IDBA

**Availability:** Acknowledgement requested

**Website Status:** Last checked up

**Abbreviations:** IDBA-UD

**Resource Name:** IDBA-UD

**Resource ID:** SCR_011912

**Alternate IDs:** OMICS_01423

Ratings and Alerts
No rating or validation information has been found for IDBA-UD.

No alerts have been found for IDBA-UD.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 224 mentions in open access literature.

Listed below are recent publications. The full list is available at FDI Lab - SciCrunch.org.


Danczak RE, et al. (2020) Ecological Assembly Processes Are Coordinated between Bacterial and Viral Communities in Fractured Shale Ecosystems. mSystems, 5(2).


Mastor NN, et al. (2020) Whole genome sequencing data of a clinical strain EGR748 from Sabah, Malaysia. Data in brief, 33, 106370.


