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/ida_u_d/

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

Resource Name: IDBA-UD

Proper Citation: IDBA-UD (RRID:SCR_011912)

Resource Type: Resource, software resource

Resource ID: SCR_011912

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

Related resources: Meta-IDBA

References: PMID: 22495754

Availability: Acknowledgement requested

Website Status: Last checked up

Alternate IDs: OMICS_01423

Abbreviations: IDBA-UD

Mentions Count: 158

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

No alerts have been found for IDBA-UD.

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**Data and Source Information**

**Source:** SciCrunch Registry

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**Usage and Citation Metrics**

We found 158 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [FDI Lab - SciCrunch Infrastructure](https://fdilab.scicrunch.org).


Göller PC, et al. (2020) Uncovering a hidden diversity: optimized protocols for the extraction of dsDNA bacteriophages from soil. Microbiome, 8(1), 17.


Seah BKB, et al. (2019) Sulfur-Oxidizing Symbionts without Canonical Genes for Autotrophic COFixation. mBio, 10(3).


