

Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 2026

EBARDenovo

RRID:SCR_011890

Type: Tool

Proper Citation

EBARDenovo (RRID:SCR_011890)

Resource Information

URL: <http://ebardenovo.sourceforge.net/>

Proper Citation: EBARDenovo (RRID:SCR_011890)

Description: Highly accurate de novo assembly of RNA-Seq with efficient chimera-detection.

Abbreviations: EBARDenovo

Resource Type: software resource

Resource Name: EBARDenovo

Resource ID: SCR_011890

Alternate IDs: OMICS_01317

Record Creation Time: 20220129T080307+0000

Record Last Update: 20260725T190722+0000

Ratings and Alerts

No rating or validation information has been found for EBARDenovo.

No alerts have been found for EBARDenovo.

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 [nidm-terms](#) .

Fisher TW, et al. (2023) Sub-optimal temperatures lead to altered expression of stress-related genes and increased 'Candidatus Liberibacter solanacearum' accumulation in potato psyllid. *Frontiers in insect science*, 3, 1279365.

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