Resource Summary Report

Generated by NIF on Apr 20, 2025

GARM

RRID:SCR_006731
Type: Tool

Proper Citation

GARM (RRID:SCR_006731)

Resource Information

URL: http://garm-meta-assem.sourceforge.net/

Proper Citation: GARM (RRID:SCR_006731)

Description: A new software pipeline to merge and reconcile assemblies from different algorithms or sequencing technologies.

Abbreviations: GARM

Synonyms: Genome Assembler Reconcilation and Merging

Resource Type: software resource

Funding:

Resource Name: GARM

Resource ID: SCR_006731

Alternate IDs: OMICS_01420

Record Creation Time: 20220129T080237+0000

Record Last Update: 20250420T014344+0000

Ratings and Alerts

No rating or validation information has been found for GARM.

No alerts have been found for GARM.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 11 mentions in open access literature.

Listed below are recent publications. The full list is available at NIF.

Hayden NK, et al. (2023) Reducing the risk of criminal exploitation using multi-systemic therapy (the RESET Study): study protocol for a feasibility study and process evaluation. Pilot and feasibility studies, 9(1), 193.

Gupta A, et al. (2021) Evaluation of immediate and short-term efficacy of DualStim therapy with and without intracavernosal umbilical cord-derived Wharton's jelly in patients with erectile dysfunction: Study protocol for a randomized controlled trial. Contemporary clinical trials communications, 23, 100790.

Vong K, et al. (2021) Disrupting tumor onset and growth via selective cell tagging (SeCT) therapy. Science advances, 7(17).

Kim J, et al. (2020) Whole-genome, transcriptome, and methylome analyses provide insights into the evolution of platycoside biosynthesis in Platycodon grandiflorus, a medicinal plant. Horticulture research, 7, 112.

Nielsen ES, et al. (2020) Multi-model seascape genomics identifies distinct environmental drivers of selection among sympatric marine species. BMC evolutionary biology, 20(1), 121.

Nicholson TL, et al. (2020) Comparative Virulence and Genomic Analysis of Streptococcus suis Isolates. Frontiers in microbiology, 11, 620843.

Gokul EA, et al. (2019) Remotely sensing harmful algal blooms in the Red Sea. PloS one, 14(4), e0215463.

de Souza WM, et al. (2019) Pingu virus: A new picornavirus in penguins from Antarctica. Virus evolution, 5(2), vez047.

Solórzano S, et al. (2019) De Novo Assembly Discovered Novel Structures in Genome of Plastids and Revealed Divergent Inverted Repeats in Mammillaria (Cactaceae, Caryophyllales). Plants (Basel, Switzerland), 8(10).

Modha S, et al. (2019) Metaviromics Reveals Unknown Viral Diversity in the Biting Midge Culicoides impunctatus. Viruses, 11(9).

Leijs R, et al. (2015) The Evolution of Epigean and Stygobitic Species of Koonunga Sayce, 1907 (Syncarida: Anaspidacea) in Southern Australia, with the Description of Three New Species. PloS one, 10(8), e0134673.