Resource Summary Report

Generated by dkNET on May 19, 2025

FastGap

RRID:SCR_018974
Type: Tool

Proper Citation

FastGap (RRID:SCR_018974)

Resource Information

URL: http://www.aubot.dk/FastGap_home.htm

Proper Citation: FastGap (RRID:SCR_018974)

Description: Software tool for assembly of DNA sequence alignment files in BioEdit fasta format into NEXUS format ready for analysis in programs such as PAUP or MrBayes. Used to calibrate gap in sequence after alignment.

Synonyms: FastGap 1.2

Resource Type: software resource, software application, data processing software

Keywords: DNA sequence alignment file, file assembly, BioEdit fasta, NEXUS format, sequence gap calibration, gap

Funding:

Availability: Free, Available for download, Freely available

Resource Name: FastGap

Resource ID: SCR_018974

Record Creation Time: 20220129T080342+0000

Record Last Update: 20250513T062030+0000

Ratings and Alerts

No rating or validation information has been found for FastGap.

No alerts have been found for FastGap.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 7 mentions in open access literature.

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

Pita S, et al. (2022) Multidisciplinary approach detects speciation within the kissing bug Panstrongylus rufotuberculatus populations (Hemiptera, Heteroptera, Reduviidae). Memorias do Instituto Oswaldo Cruz, 116, e210259.

Fehrer J, et al. (2022) A Multigene Phylogeny of Native American Hawkweeds (Hieracium Subgen. Chionoracium, Cichorieae, Asteraceae): Origin, Speciation Patterns, and Migration Routes. Plants (Basel, Switzerland), 11(19).

Lumini E, et al. (2020) Native Arbuscular Mycorrhizal Fungi Characterization from Saline Lands in Arid Oases, Northwest China. Journal of fungi (Basel, Switzerland), 6(2).

de Rezende Dias G, et al. (2018) Cryptic diversity in an Atlantic Forest malaria vector from the mountains of South-East Brazil. Parasites & vectors, 11(1), 36.

Cheng J, et al. (2017) Cryptic diversity in the Japanese mantis shrimp Oratosquilla oratoria (Crustacea: Squillidae): Allopatric diversification, secondary contact and hybridization. Scientific reports, 7(1), 1972.

Grace OM, et al. (2015) Evolutionary history and leaf succulence as explanations for medicinal use in aloes and the global popularity of Aloe vera. BMC evolutionary biology, 15, 29.

Wang Z, et al. (2014) Phylogeny reconstruction and hybrid analysis of populus (Salicaceae) based on nucleotide sequences of multiple single-copy nuclear genes and plastid fragments. PloS one, 9(8), e103645.