Resource Summary Report

Generated by NIF on Apr 17, 2025

Easyfig

RRID:SCR_013169

Type: Tool

Proper Citation

Easyfig (RRID:SCR_013169)

Resource Information

URL: http://easyfig.sourceforge.net/

Proper Citation: Easyfig (RRID:SCR_013169)

Description: A Python application for creating linear comparison figures of multiple genomic

loci with an easy-to-use graphical user interface (GUI).

Abbreviations: Easyfig

Resource Type: software resource

Funding:

Resource Name: Easyfig

Resource ID: SCR_013169

Alternate IDs: OMICS_00907

Record Creation Time: 20220129T080314+0000

Record Last Update: 20250410T070339+0000

Ratings and Alerts

No rating or validation information has been found for Easyfig.

No alerts have been found for Easyfig.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 953 mentions in open access literature.

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

Almeida-Santos AC, et al. (2025) The healthy human gut can take it all: vancomycin-variable, linezolid-resistant strains and specific bacteriocin-species interplay in Enterococcus spp. Applied and environmental microbiology, 91(1), e0169924.

Latorre-Fernández J, et al. (2025) Evaluation of the double-zone hemolysis (DZH) test for the detection of livestock-associated methicillin-resistant Staphylococcus aureus. Microbiology spectrum, 13(1), e0110224.

Long D, et al. (2025) Epidemiological and genetic charateristics of Vibrio vulnificus from diverse sources in China during 2012-2023. Communications biology, 8(1), 9.

Zamunér CFC, et al. (2025) Evolution and spread of Xanthomonas citri subsp. citri in the São Paulo, Brazil, citrus belt inferred from 758 novel genomes. Microbial genomics, 11(1).

Larsen AL, et al. (2025) Hospital toilets and drainage systems as a reservoir for a long-term polyclonal outbreak of clinical infections with multidrug-resistant Klebsiella oxytoca species complex. Infection prevention in practice, 7(1), 100430.

Kokoulin MS, et al. (2025) Structure of a Sulfated Capsular Polysaccharide from the Marine Bacterium Cobetia marina KMM 1449 and a Genomic Insight into Its Biosynthesis. Marine drugs, 23(1).

Xie Y, et al. (2025) The prevalence of optrA-carrying Enterococci in the vaginal micro-ecology of pregnant women in late pregnancy. Microbiology spectrum, 13(1), e0213524.

Quesille-Villalobos AM, et al. (2025) Multispecies emergence of dual blaKPC/NDM carbapenemase-producing Enterobacterales recovered from invasive infections in Chile. Antimicrobial agents and chemotherapy, 69(1), e0120524.

He Z, et al. (2025) Characterization and genomic analysis of the highly virulent Acinetobacter baumannii ST1791 strain dominating in Anhui, China. Antimicrobial agents and chemotherapy, 69(1), e0126224.

Yehia FAA, et al. (2025) From Isolation to Application: Utilising Phage-Antibiotic Synergy in Murine Bacteremia Model to Combat Multidrug-Resistant Enterococcus faecalis. Microbial biotechnology, 18(1), e70075.

Hsu TK, et al. (2025) Characterization and genome analysis of a novel phage BP15 infecting

Vibrio parahaemolyticus. Scientific reports, 15(1), 2801.

Banar M, et al. (2025) A novel broad-spectrum bacteriophage cocktail against methicillinresistant Staphylococcus aureus: Isolation, characterization, and therapeutic potential in a mastitis mouse model. PloS one, 20(1), e0316157.

Puangseree J, et al. (2024) Molecular basis of the persistence of chloramphenicol resistance among Escherichia coli and Salmonella spp. from pigs, pork and humans in Thailand. PloS one, 19(5), e0304250.

Abdullahi IN, et al. (2024) Genomic Characterization and Phylogenetic Analysis of Linezolid-Resistant Enterococcus from the Nostrils of Healthy Hosts Identifies Zoonotic Transmission. Current microbiology, 81(8), 225.

Chen J, et al. (2024) Genotypic and phenotypic characteristics of Acinetobacter baumannii isolates from the people's hospital of Qingyang City, Gansu province. BMC genomics, 25(1), 727.

Hatrongjit R, et al. (2024) Genomic analysis of carbapenem- and colistin-resistant Klebsiella pneumoniae complex harbouring mcr-8 and mcr-9 from individuals in Thailand. Scientific reports, 14(1), 16836.

Wang W, et al. (2024) Whole genome sequencing insight into carbapenem-resistant and multidrug-resistant Acinetobacter baumannii harboring chromosome-borne blaOXA-23. Microbiology spectrum, 12(9), e0050124.

Chen HX, et al. (2024) Two colistin resistance-producing Aeromonas strains, isolated from coastal waters in Zhejiang, China: characteristics, multi-drug resistance and pathogenicity. Frontiers in microbiology, 15, 1401802.

Nair S, et al. (2024) Presence of phage-plasmids in multiple serovars of Salmonella enterica. Microbial genomics, 10(5).

Shen W, et al. (2024) Mapping the widespread distribution and transmission dynamics of linezolid resistance in humans, animals, and the environment. Microbiome, 12(1), 52.