

Resource Summary Report

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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

Resource Name: Easyfig

Resource ID: SCR_013169

Alternate IDs: OMICS_00907

Record Creation Time: 20220129T080314+0000

Record Last Update: 20260801T190456+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 1101 mentions in open access literature.

Listed below are recent publications. The full list is available at [ASWG](#) .

Tian R, et al. (2026) Emergence and genomic characterization of mcr-8.2-positive *Klebsiella pneumoniae* from layer chickens in Xinjiang, China. *Poultry science*, 105(1), 106190.

Figuerola Y, et al. (2026) STEC in the natural environment of Uruguay: genomic surveillance and environmental circulation in the framework of One Health. *Microbiology spectrum*, 14(1), e0315325.

Yeki S, et al. (2026) Sex-Specific doublesex Regulation Targeting the Color-Patterning Gene *h* Underlies the Evolution of Wing Sexual Dimorphism in the Harlequin Ladybug *Harmonia axyridis*. *Evolution & development*, 28(1), e70028.

Hu Y, et al. (2026) Global epidemiological and genetic characteristics of carbapenem-resistant *Escherichia coli* carrying blaIMP. *Microbiology spectrum*, 14(1), e0324425.

Sia CM, et al. (2025) Distinct adaptation and epidemiological success of different genotypes within *Salmonella enterica* serovar Dublin. *eLife*, 13.

Biedrzycka M, et al. (2025) Carbapenemase-Producing Enterobacterales from Patients Arriving from Ukraine in Poland, March 2022-February 2023. *Infectious diseases and therapy*, 14(2), 401.

Jiang H, et al. (2025) Prevalence and characterization of class I integrons in multidrug-resistant *Escherichia coli* isolates from humans and food-producing animals in Zhejiang Province, China. *BMC microbiology*, 25(1), 76.

Fujii A, et al. (2025) Comprehensive analysis of bacteriocins produced by clinical enterococcal isolates and their antibacterial activity against Enterococci including VRE. *Scientific reports*, 15(1), 4846.

Liu H, et al. (2025) Characteristics of the smallest brucellaphage with strong lytic ability. *Frontiers in veterinary science*, 12, 1530123.

Krutova M, et al. (2025) Genomic islands and molecular mechanisms relating to drug-resistance in *Clostridioides (Clostridium) difficile* PCR ribotype 176. *Emerging microbes & infections*, 14(1), 2482698.

Feng Y, et al. (2025) Identification and evolution of novel cfr-carrying plasmids in XDR *Klebsiella pneumoniae* strains from a chicken farm. *Microbiology spectrum*, 13(5), e0262824.

Howard-Varona C, et al. (2025) Infection and Genomic Properties of Single- and Double-Stranded DNA Cellulophaga Phages. *Viruses*, 17(3).

Fu S-Y, et al. (2025) Optimizing phage therapy for carbapenem-resistant *Enterobacter cloacae* bacteremia: insights into dose and timing. *Antimicrobial agents and chemotherapy*, 69(4), e0168324.

Kyrkou I, et al. (2025) *Pseudomonas aeruginosa* maintains an inducible array of novel and diverse prophages over lengthy persistence in cystic fibrosis lungs. *FEMS microbiology letters*, 372.

Chai J, et al. (2025) Isolation, characterization, and application of the novel polyvalent bacteriophage vB_EcoM_XAM237 against pathogenic *Escherichia coli*. *Veterinary research*, 56(1), 90.

Becerra-Aparicio F, et al. (2025) Whole Genome Sequencing Analysis of *Klebsiella pneumoniae* Isolates from Health Care-Associated Bacteremia of Urinary Origin in Spain: Findings from the Multicenter ITUBRAS-2 Cohort Study. *Open forum infectious diseases*, 12(4), ofaf164.

Xu L, et al. (2025) Genomic Insights into the Pathogenicity of Hypervirulent *Aeromonas hydrophila* Strain D4 Isolated from Diseased Blunt Snout Bream with the Epidemic Sequence Type 251 Clones. *Pathogens (Basel, Switzerland)*, 14(6).

Liu JQ, et al. (2025) Genome-Based Reclassification of Two *Haloarcula* Species and Characterization of *Haloarcula montana* sp. nov. *Biology*, 14(6).

Wu X, et al. (2025) A Novel Metallo- β -Lactamase AMM-1 From *Alteromonas mangrovi* Reveals a Cryptic Environmental Reservoir of Carbapenem Resistance. *Microbial biotechnology*, 18(7), e70191.

Kim EJ, et al. (2025) Bacteriophage cocktail LEC2-LEC10 for broad-spectrum control of pathogenic and uncharacterized *Escherichia coli* in fresh produce. *Frontiers in microbiology*, 16, 1594533.