

Resource Summary Report

Generated by Kravitz on Sep 18, 2026

FinchTV

RRID:SCR_005584

Type: Tool

Proper Citation

FinchTV (RRID:SCR_005584)

Resource Information

URL: <http://www.geospiza.com/Products/finchtv.shtml>

Proper Citation: FinchTV (RRID:SCR_005584)

Description: Chromatogram viewer that can display an entire trace in a scalable multi-pane view, allows raw data views, BLAST searching and the ability to reverse complement sequences and traces., THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 16,2025.

Abbreviations: FinchTV

Synonyms: FinchTV - A Brilliant Trace Viewer

Resource Type: commercial organization, software resource

Keywords: chromatogram, linux, mac osx, windows, solaris, dna sequence trace, dna sequence, trace

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: FinchTV

Resource ID: SCR_005584

Alternate IDs: OMICS_01019

Record Creation Time: 20220129T080231+0000

Record Last Update: 20260912T200315+0000

Ratings and Alerts

No rating or validation information has been found for FinchTV.

No alerts have been found for FinchTV.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 646 mentions in open access literature.

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

Phaka FM, et al. (2025) Barcoding and traditional health practitioner perspectives are informative to monitor and conserve frogs and reptiles traded for traditional medicine in urban South Africa. *Molecular ecology resources*, 25(2), e13873.

Legnardi M, et al. (2025) The role of the rural poultry sector in viral circulation and evolution: evaluation of infectious bronchitis virus and avian metapneumovirus dynamics in an Italian dealer farm. *Poultry science*, 104(9), 105420.

Dinh QM, et al. (2025) Genetic Relationship and Diversity of *Ellochelon vaigiensis* in the Vietnamese Mekong Delta Using Cytochrome b Gene Sequences. *Ecology and evolution*, 15(6), e71639.

Benigno A, et al. (2025) Morphological, Molecular and Pathological Characterization of *Phytophthora pseudocryptogea* Associated with *Rosmarinus officinalis* Dieback in Tuscany, Central Italy. *Microorganisms*, 13(3).

Winiarczyk D, et al. (2025) A simple validation and screening method for CRISPR/Cas9-mediated gene editing in mouse embryos to facilitate genetically modified mice production. *PloS one*, 20(3), e0312722.

Magagula M, et al. (2025) Investigating the antimicrobial and anticancer potential of culturable fungal endophytes isolated from the stems of *Kirkia acuminata* Oliv. *BMC microbiology*, 25(1), 343.

Gunyakti Kilinc S, et al. (2025) A preliminary study on helminths detected in red foxes (*Vulpes vulpes*) in Bingol Province of Türkiye: morphological and molecular approaches. *BMC veterinary research*, 21(1), 391.

Dudaš T, et al. (2025) Diversity and Patulin Production of *Penicillium* spp. Associated with Apple Blue Mold in Serbia. *Journal of fungi (Basel, Switzerland)*, 11(3).

Wetters S, et al. (2025) B-class gene GLOBOSA - a facilitator for enriched species diversity of *Salvia* in the New World? *Plant biology (Stuttgart, Germany)*, 27(3), 333.

Tayar S, et al. (2025) *Ganoderma lucidum* Immobilized on Wood Demonstrates High Persistence During the Removal of OPFRs in a Trickle-Bed Bioreactor. *Journal of fungi (Basel, Switzerland)*, 11(2).

Dorszewska J, et al. (2025) Genetic variants of ZNF746 and the level of plasma Parkin, PINK1, and ZNF746 proteins in patients with Parkinson's disease. *IBRO neuroscience reports*, 18, 342.

Sumaira S, et al. (2025) Molecular epidemiology and phylogeny of *Anaplasma* species in goats from Pakistan. *PloS one*, 20(5), e0325467.

Olaitan MO, et al. (2025) Gender-specific antibiotic-resistance patterns and characterization of bacterial flora of undergraduate students' mobile phones and associated factors at a Nigerian university. *BMC microbiology*, 25(1), 30.

Dercksen M, et al. (2025) Diagnosis of Primary Trimethylaminuria in an Affected Patient With a Rare Genotype in Sub-Saharan Africa. *JIMD reports*, 66(2), e70005.

Mvogo Nyebe RA, et al. (2025) Multi-omics characterization of aflatoxigenic *Aspergillus* from grains and rhizosphere of maize across agroecological zones of Cameroon. *Scientific reports*, 15(1), 18407.

Shabbir R, et al. (2025) Leading report on the molecular prevalence of emerging pathogens *Hepatozoon* sp. and *Lankesterella* sp. in the blood samples of seven wild lizard species. *Scientific reports*, 15(1), 9014.

Di Cristina G, et al. (2025) Pioneering genome editing in parthenogenetic stick insects: CRISPR/Cas9-mediated gene knockout in *Medauroidea extradentata*. *Scientific reports*, 15(1), 2584.

Kumar N, et al. (2025) Impact of altitudinal variations on plant growth dynamics, nutritional composition, and free living rhizospheric N₂ fixing bacterial community of *Eruca sativa*. *Scientific reports*, 15(1), 13839.

Dami FD, et al. (2025) The Eye of the Chiropterologist: Phenotypic Versus Genotypic Identification of Bats. *Ecology and evolution*, 15(6), e71561.

Angel-Ruiz NA, et al. (2025) Bioprospecting of four *Beauveria bassiana* strains and their potential as biological control agents for *Anastrepha ludens* Loew 1873 (Diptera: Tephritidae). *PloS one*, 20(6), e0324441.