

Resource Summary Report

Generated by [kravitz2](#) 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 [kravitz2](#) .

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.

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.

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).

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.

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.

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.

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.

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.

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).

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.

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.

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.