# **Resource Summary Report**

Generated by dkNET on Apr 30, 2025

# Geospiza

RRID:SCR\_003264

Type: Tool

## **Proper Citation**

Geospiza (RRID:SCR\_003264)

#### **Resource Information**

URL: http://www.geospiza.com/

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**Description:** Developer of enterprise-class software systems for workflow management of genetic analysis. Geospiza has established itself as the life science industry's top software developer for meeting the combined laboratory, data management and analytical needs of biotechnology and pharmaceutical companies, universities, researchers, contract core and diagnostic laboratories involved in genetic testing and manufacturing bio-therapeutics. Geospiza's products are designed and built by biologists, enabling researchers to focus on the science, not the software. \*GeneSifter Analysis Edition: Access easy-to-use statistical, visualization and annotation tools for Microarray and Next Generation Sequencing data. \*GeneSifter Lab Edition: Manage cross-platform core lab operations from a single interface out-of-the-box.

Abbreviations: Geospiza

Synonyms: Geospiza Inc.

**Resource Type:** software resource, material service resource, service resource, commercial organization, production service resource

Keywords: genetic analysis, microarray, next generation sequencing, data management

Funding:

Availability: Commercial license

Resource Name: Geospiza

Resource ID: SCR\_003264

Alternate IDs: OMICS\_01127, nif-0000-31431

Old URLs: http://www.geospiza.com/finchtv.html

**Record Creation Time:** 20220129T080218+0000

**Record Last Update:** 20250430T055212+0000

## **Ratings and Alerts**

No rating or validation information has been found for Geospiza.

No alerts have been found for Geospiza.

#### **Data and Source Information**

Source: SciCrunch Registry

## **Usage and Citation Metrics**

We found 103 mentions in open access literature.

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

Sonowal H, et al. (2025) Preclinical Development of Tuspetinib for the Treatment of Acute Myeloid Leukemia. Cancer research communications, 5(1), 74.

Che Alias MA, et al. (2024) Morphological and genetic identification of Halophila species and a new distribution record of Halophila nipponica at the Tanjung Adang Laut shoal, Johor, Malaysia. PloS one, 19(10), e0309143.

Karakoc S, et al. (2024) Genetic diversity and haplotypes of Cysticercus tenuicollis isolates from slaughtered sheep and goats in Elazig and Bingol provinces of Turkey. Veterinary medicine and science, 10(4), e1411.

Jaiswal P, et al. (2024) Morphological and Molecular Characterization of Trichoderma Isolates from Vegetable Crop Rhizospheres in Nepal. F1000Research, 13, 1088.

Lea V, et al. (2024) Toward the Development of the Trojan Female Technique in Pest Insects: Male-Specific Influence of Mitochondrial Haplotype on Reproductive Output in the Seed Beetle Acanthoscelides obtectus. Evolutionary applications, 17(12), e70065.

Kisielinski K, et al. (2024) The bacterial burden of worn face masks-observational research and literature review. Frontiers in public health, 12, 1460981.

Pena R, et al. (2024) Ancestry of the major long-range regulatory site of the ?-globin genes in the Portuguese population with the common 3.7 kb ?-thalassemia deletion. Molecular biology reports, 51(1), 612.

Ahasan SA, et al. (2023) Wide variation of heterozygotic genotypes of recent fasciolid hybrids from livestock in Bangladesh assessed by rDNA internal transcribed spacer region sequencing and cloning. One health (Amsterdam, Netherlands), 17, 100614.

Augusto DG, et al. (2023) A common allele of HLA is associated with asymptomatic SARS-CoV-2 infection. Nature, 620(7972), 128.

Novak Babi? M, et al. (2023) Effect of Location, Disinfection, and Building Materials on the Presence and Richness of Culturable Mycobiota through Oligotrophic Drinking Water Systems. Journal of fungi (Basel, Switzerland), 9(11).

Tegegne D, et al. (2023) First genetic detection and characterization of canine parvovirus type 2 (Carnivore protoparvovirus 1) in southwestern Ethiopia. Veterinary research communications, 47(2), 975.

Grassi A, et al. (2023) A Lethal Case of Disseminated Cladosporium allicinum Infection in a Captive African Bullfrog. Journal of fungi (Basel, Switzerland), 9(2).

Kosovac A, et al. (2023) Epidemiological role of novel and already known 'Ca. P. solani' cixiid vectors in rubbery taproot disease of sugar beet in Serbia. Scientific reports, 13(1), 1433.

Dougue AN, et al. (2022) Multilocus sequence typing (MLST) analysis reveals many novel genotypes and a high level of genetic diversity in Candida tropicalis isolates from Italy and Africa. Mycoses, 65(11), 989.

de Lillo E, et al. (2022) Integrative Taxonomy and Synonymization of Aculus mosoniensis (Acari: Eriophyidae), a Potential Biological Control Agent for Tree of Heaven (Ailanthus altissima). Insects, 13(5).

Kneubehl AR, et al. (2022) Amplification and sequencing of entire tick mitochondrial genomes for a phylogenomic analysis. Scientific reports, 12(1), 19310.

Franzo G, et al. (2021) Canine Circovirus in Foxes from Northern Italy: Where Did It All Begin? Pathogens (Basel, Switzerland), 10(8).

Novak Babi? M, et al. (2021) Water-Transmitted Fungi are Involved in Degradation of Concrete Drinking Water Storage Tanks. Microorganisms, 9(1).

Xu J, et al. (2021) An Axin2 mutation and perinatal risk factors contribute to sagittal craniosynostosis: evidence from a Chinese female monochorionic diamniotic twin family. Hereditas, 158(1), 20.

Common LK, et al. (2021) Avian vampire fly (Philornis downsi) mortality differs across Darwin's finch host species. Scientific reports, 11(1), 15832.