Resource Summary Report

Generated by Kravitz on Sep 10, 2026

Golden Helix Incorporated

RRID:SCR_012191
Type: Tool

Proper Citation

Golden Helix Incorporated (RRID:SCR_012191)

Resource Information

URL: http://goldenhelix.com/

Proper Citation: Golden Helix Incorporated (RRID:SCR_012191)

Description: Specializes in sequence and array-based SNP and copy number analysis, genetic association software, and analytic services. Their technologies empower scientists to determine the genetic causes of disease, transform drug discovery, develop genetic diagnostics, and advance the quest for personalized medicine.

Abbreviations: Golden Helix

Synonyms: Golden Helix Inc., GoldenHelix.com

Resource Type: commercial organization

Keywords: resource, portal, analysis, software

Availability: Available to external user

Resource Name: Golden Helix Incorporated

Resource ID: SCR_012191

Alternate IDs: SciEx_10349

Old URLs: http://www.goldenhelix.com/Services, http://www.scienceexchange.com/facilities/golden-helix-inc

Record Creation Time: 20220129T080308+0000

Record Last Update: 20260905T132722+0000

Ratings and Alerts

No rating or validation information has been found for Golden Helix Incorporated.

No alerts have been found for Golden Helix Incorporated.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 170 mentions in open access literature.

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

Tom W, et al. (2026) A Whole-Exome Sequencing-Based Exploration of Chronic Kidney Disease of Unknown Etiology (CKDu) in an Endemic Population in Sri Lanka. International journal of molecular sciences, 27(8).

Ricci C, et al. (2026) Adult granulosa cell tumours of the testis analogous to ovarian counterparts are exceptionally rare: analysis of a multicentric series and review of the literature. Histopathology, 88(4), 831.

Zhang S, et al. (2026) Genome-Wide Association Study Suggests rrp44 Is a Key Regulator of Growth Traits in Channel Catfish (Ictalurus punctatus). Current issues in molecular biology, 48(4).

Côrtes L, et al. (2026) Family-Based Study Reveals PDE11A/PDE11A-AS1 Variants in Testicular Germ Cell Tumor Predisposition. International journal of molecular sciences, 27(12).

Liu D, et al. (2025) Assessing SARS-CoV-2 Rare Mutations and Transmission in New York City by NGS. Microorganisms, 13(8).

Olou AA, et al. (2025) EV DNA from pancreatic cancer patient-derived cells harbors molecular, coding, non-coding signatures and mutational hotspots. Communications biology, 8(1), 368.

Kristiansen MH, et al. (2025) Prevalence and impact of additional CHIP mutations in JAK2V617F-positive ischaemic cerebrovascular patients. British journal of haematology, 207(3), 1029.

Thorsrud JA, et al. (2025) Breeding values and index creation for health and behavior traits in Labrador Retriever guide dogs. Frontiers in veterinary science, 12, 1628161.

Thorsrud JA, et al. (2025) Performance Comparison of Genomic Best Linear Unbiased Prediction and Four Machine Learning Models for Estimating Genomic Breeding Values in Working Dogs. Animals : an open access journal from MDPI, 15(3).

Yang Z, et al. (2025) Sequencing validates deep learning models for EHR-based detection of Noonan syndrome in pediatric patients. NPJ genomic medicine, 10(1), 56.

Icedo-Nuñez S, et al. (2025) Validation of Polymorphisms Associated with the Immune Response After Vaccination Against Porcine Reproductive and Respiratory Syndrome Virus in Yorkshire Gilts. Veterinary sciences, 12(4).

Beaman GM, et al. (2025) Case Report: Prolonged survival in Schinzel-Giedion syndrome featuring megaureter and de novo SETBP1 mutation. Frontiers in pediatrics, 13, 1534192.

Ali GS, et al. (2025) GWAS identifies a polyembryony locus in mango: development of KASP and PACE markers for marker-assisted breeding. Frontiers in plant science, 16, 1508027.

Reshadmanesh A, et al. (2024) First Case of Macrocephaly, Dysmorphic Facies, and Psychomotor Retardation Harboring Co-inherited Variants in HERC1 and PMP22 Genes from Iran: Two Novel Variants. Archives of Iranian medicine, 27(12), 700.

Nisa WU, et al. (2024) Insights into maydis leaf blight resistance in maize: a comprehensive genome-wide association study in sub-tropics of India. BMC genomics, 25(1), 760.

Glenthøj A, et al. (2024) DAHEAN: A Danish nationwide study ensuring quality assurance through real-world data for suspected hereditary anemia patients. Orphanet journal of rare diseases, 19(1), 284.

Contreras-Méndez LA, et al. (2024) The Anti-Müllerian Hormone as Endocrine and Molecular Marker Associated with Reproductive Performance in Holstein Dairy Cows Exposed to Heat Stress. Animals : an open access journal from MDPI, 14(2).

Zorc M, et al. (2024) Selection Signatures Reveal Candidate Genes for the Cornish Rex Breed-Specific Phenotype. Genes, 15(3).

Thompson MA, et al. (2024) Population genomics provide insight into ancestral relationships and diversity of the feral horses of Theodore Roosevelt National Park. Ecology and evolution, 14(4), e11197.

Šimon M, et al. (2024) Whole genome sequencing of mouse lines divergently selected for fatness (FLI) and leanness (FHI) revealed several genetic variants as candidates for novel obesity genes. Genes & genomics, 46(5), 557.