

Resource Summary Report

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Structure Harvester

RRID:SCR_017636

Type: Tool

Proper Citation

Structure Harvester (RRID:SCR_017636)

Resource Information

URL: <http://taylor0.biology.ucla.edu/structureHarvester/>

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Description: Web based program for collating results generated by program STRUCTURE. Provides assess and visualize likelihood values across multiple values of K and hundreds of iterations for easier detection of number of genetic groups that best fit data. Reformats data for use in downstream programs, such as CLUMPP. It is complement for using software Structure in genetics population. Website and program for visualizing STRUCTURE output and implementing Evanno method., THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 16, 2025.

Synonyms: StructureHarvester

Resource Type: web service, software resource, production service resource, data access protocol, service resource, analysis service resource

Defining Citation: [DOI:10.1007/s12686-011-9548-7](https://doi.org/10.1007/s12686-011-9548-7)

Keywords: Visualizing, STRUCTURE, Evanno, method, collating, result, detection, genetic, group, fit, data, reformat

Funding: NCI U24 CA143858;
NCI R21 CA135937

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: Structure Harvester

Resource ID: SCR_017636

Record Creation Time: 20220129T080336+0000

Record Last Update: 20260803T040736+0000

Ratings and Alerts

No rating or validation information has been found for Structure Harvester.

No alerts have been found for Structure Harvester.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 297 mentions in open access literature.

Listed below are recent publications. The full list is available at [kravitz2](#) .

Kniesz K, et al. (2025) High genomic connectivity within Anatoma at hydrothermal vents along the Central and Southeast Indian Ridge. Scientific reports, 15(1), 1971.

Teklewold TH, et al. (2025) Genetic diversity and population structure of Ethiopian Sweet Sorghum [*Sorghum bicolor* (L.) Moench] germplasms using SSR markers. PloS one, 20(3), e0316549.

Belkina D, et al. (2025) In-depth population genetic study of *Vitis vinifera* ssp. *sylvestris* from the Black Sea region and its virome. Frontiers in plant science, 16, 1536862.

Patel MK, et al. (2025) Genome-wide association study uncovers key genomic regions governing agro-morphological and quality traits in Indian mustard [*Brassica juncea* (L.) Czern. and Coss.]. PloS one, 20(4), e0322120.

Ruppert LS, et al. (2025) Gene Flow Across Large Distances in the Cavity-Nesting Wasp *Deuteragenia subintermedia* in a Central European Forest. Ecology and evolution, 15(4), e71294.

Toker TP, et al. (2025) Genomic insights into Mediterranean pepper diversity using ddRADSeq. PloS one, 20(3), e0318105.

Rakshith HS, et al. (2025) Genetic diversity of the zigzag ladybird beetle, *Cheilomenes sexmaculata* (F.) (Coleoptera: Coccinellidae) with its distribution in India and implications for biological control. Scientific reports, 15(1), 15629.

Taboada S, et al. (2025) Connectivity and Adaptation Patterns of the Deep-Sea Ground-Forming Sponge *Geodia hentscheli* Across Its Entire Distribution. Molecular biology and evolution, 42(7).

Gyawali S, et al. (2025) Genome wide association study of Fusarium wilt resistance in *Spinacia turkestanica*. Scientific reports, 15(1), 15486.

Ma X, et al. (2025) ddRAD-seq data reveals the significant population structure and potential causes of population divergence in *Bougainvillea*. Scientific reports, 15(1),

34200.

Wang Q, et al. (2025) Population Patterns and Dynamics of *Ilisha elongata* (Clupeiformes: Pristigasteridae) Revealed by Target Enrichment Data. *Evolutionary applications*, 18(8), e70142.

Li Y, et al. (2025) Migration of wheat stripe rust from the primary overwintering region to neighboring regions in China. *Communications biology*, 8(1), 350.

Yang H, et al. (2025) Population Genetics of *Haliotis discus hannai* in China Inferred Through EST-SSR Markers. *Genes*, 16(1).

Bernád V, et al. (2024) Unlocking the genetic diversity and population structure of the newly introduced two-row spring European Heritage Barley collection (ExHIBiT). *Frontiers in plant science*, 15, 1268847.

Arévalo-Marín E, et al. (2024) Genetic analyses and dispersal patterns unveil the Amazonian origin of guava domestication. *Scientific reports*, 14(1), 15755.

de Sales RP, et al. (2024) Addressing Conservation Needs: Genetic Diversity and Population Ecology of the Endemic Tree *Spondias tuberosa* Arruda. *Scientifica*, 2024, 5023974.

Faria JCT, et al. (2024) Genetic resources of African mahogany in Brazil: genomic diversity and structure of forest plantations. *BMC plant biology*, 24(1), 858.

Zheng JX, et al. (2024) Species Composition of a Small Mammal Community and Prevalence of *Echinococcus* spp. in the Alpine Pastoral Area of the Eastern Tibetan Plateau. *Pathogens (Basel, Switzerland)*, 13(7).

Meulenbroek P, et al. (2024) Small-scale metapopulation structure of a limnophilic fish species in a natural river system investigated using microsatellite genotyping by amplicon sequencing (SSR-GBAS). *BMC ecology and evolution*, 24(1), 1.

Sallam A, et al. (2024) Genome-wide analysis for root and leaf architecture traits associated with drought tolerance at the seedling stage in a highly ecologically diverse wheat population. *Computational and structural biotechnology journal*, 23, 870.