

Resource Summary Report

Generated by [PRECISE-TBI](#) on Aug 1, 2026

Genoscope

RRID:SCR_002172

Type: Tool

Proper Citation

Genoscope (RRID:SCR_002172)

Resource Information

URL: <http://www.genoscope.cns.fr/spip/spip.php?lang=en>

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Description: French national sequencing center with the following resources: * Sequencing ** Genoscope Projects * Environmental genomics ** Microbial diversity in wastewater ** Metabolic genomics * Bioinformatics ** Atelier for comparative genomics ** Computational Systems Biology ** Servers resources *** GGB for Generic Genome Browser: graphic interface for various databases (sequence, annotation, syntenies...) for a given organism. *** MaGe for Magnifying Microbial Genomes: annotation system for microbial genomes.

Abbreviations: Genoscope

Synonyms: Genoscope - Centre National de S?quen?age, Genoscope - French National Sequencing Center, French National Sequencing Center, Genoscope - Centre National de Sequencage

Resource Type: institution

Keywords: environmental genomics, biocatalysis, environment, genomics, sequencing, bioinformatics, biodiversity, blast, blat, ggb, mage, metabolic, whole genome shotgun, chromosome 3, cdna, chromosome 14, alternative splicing, o??kopleura dioica, mutation, enzymatic cloning, screening, syteny, data set, genome, sequence, annotation, genome browser, FASEB list

Availability: Free, Freely available

Resource Name: Genoscope

Resource ID: SCR_002172

Alternate IDs: ISNI: 0000 0004 0641 2997, Wikidata: Q3100800, grid.434728.e, nif-0000-20957

Alternate URLs: <https://ror.org/028pnqf58>

Record Creation Time: 20220129T080211+0000

Record Last Update: 20260725T190517+0000

Ratings and Alerts

No rating or validation information has been found for Genoscope.

No alerts have been found for Genoscope.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 158 mentions in open access literature.

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

Bigey F, et al. (2025) Insights into the genomic and phenotypic diversity of *Monosporozyma unispora* strains isolated from anthropic environments. *FEMS yeast research*, 25.

Ghiglione JF, et al. (2025) Mission Tara Microplastics: a holistic set of protocols and data resources for the field investigation of plastic pollution along the land-sea continuum in Europe. *Environmental science and pollution research international*, 32(16), 10032.

Cerqueira de Araujo A, et al. (2025) Genome sequences of four *Ixodes* species expands understanding of tick evolution. *BMC biology*, 23(1), 17.

Davis JT, et al. (2025) A chromosome-level genome assembly of the varied leaved jewelflower, *Streptanthus diversifolius*, reveals a recent whole genome duplication. *G3 (Bethesda, Md.)*, 15(4).

Rodríguez-Miret X, et al. (2025) Molecular complexity of the differential growth of freshwater diatoms along pH gradients. *ISME communications*, 5(1), ycaf078.

Tuomela H, et al. (2025) Conversion and upgrading of syringate by *Acinetobacter baylyi* ADP1. *Microbial cell factories*, 24(1), 209.

Dawson RA, et al. (2025) Carbon monoxide-oxidising *Pseudomonadota* on volcanic deposits. *Environmental microbiome*, 20(1), 12.

Tran Lu Y A, et al. (2025) Comparative Population Genomics Unveils Congruent Secondary Suture Zone in Southwest Pacific Hydrothermal Vents. *Molecular biology and evolution*, 42(2).

Dvorak E, et al. (2025) Recombination landscape and karyotypic variations revealed by linkage mapping in the grapevine downy mildew pathogen *Plasmopara viticola*. *G3* (Bethesda, Md.), 15(1).

Freyria NJ, et al. (2025) Shotgun metagenomics reveals the flexibility and diversity of Arctic marine microbiomes. *ISME communications*, 5(1), ycaf007.

Lira Garrido J, et al. (2025) The genomic history of Iberian horses since the last Ice Age. *Nature communications*, 16(1), 7098.

Pavloudi C, et al. (2025) First release of the European marine omics biodiversity observation network (EMO BON) shotgun metagenomics data from water and sediment samples. *Biodiversity data journal*, 13, e143585.

Zhou X, et al. (2025) Chromosome-level genome assembly of *Phytoseiulus persimilis* Athias-Henriot. *Scientific data*, 12(1), 293.

Elizondo-Reyna E, et al. (2025) Insights from a Genome-Wide Study of *Pantoea agglomerans* UADEC20: A Promising Strain for Phosphate Solubilization and Exopolysaccharides Production. *Current issues in molecular biology*, 47(1).

Blicharz S, et al. (2025) Laser dissection-assisted phloem transcriptomics highlights the metabolic and physiological changes accompanying clubroot disease progression in oilseed rape. *The Plant journal : for cell and molecular biology*, 121(1), e17156.

Rio S, et al. (2025) Genome-wide association for agro-morphological traits in a triploid banana population with large chromosome rearrangements. *Horticulture research*, 12(2), uhae307.

Salmon A, et al. (2025) On the way to diploidization and unexpected ploidy in the grass *Sporobolus* section *Spartina* mesopolyploids. *Nature communications*, 16(1), 1997.

Zhang Y, et al. (2025) Integrative omics analysis reveals the genetic basis of fatty acid composition in *Brassica napus* seeds. *Genome biology*, 26(1), 83.

Herbert L, et al. (2025) dCas9-SPO11-1 locally stimulates meiotic recombination in rice. *Frontiers in plant science*, 16, 1580225.

Deutschmann IM, et al. (2024) Disentangling microbial networks across pelagic zones in the tropical and subtropical global ocean. *Nature communications*, 15(1), 126.