Resource Summary Report

Generated by RRID on Aug 5, 2026

Centre dEtude du Polymorphisme Humain

RRID:SCR_008026
Type: Tool

Proper Citation

Centre dEtude du Polymorphisme Humain (RRID:SCR_008026)

Resource Information

URL: http://www.cephb.fr/

Proper Citation: Centre dEtude du Polymorphisme Humain (RRID:SCR_008026)

Description: The Centre d''Etude du Polymorphisme Humain (CEPH) is a research laboratory, the main activities of which are the setting up, storage, processing and distribution of DNA collections for the identification of genetic factors conferring susceptibility to complex disorders. These collections are established in partnership and full collaboration with external French or international research groups. The Foundation currently hosts the CEPH reference panel, the HGDP panel (Human genome Diversity Cell Line Panel) and several collections amounting mid-2008 to more than 250 000 samples. The goal of CEPH is to understand complex multifactorial disorders necessitates the establishment of structures facilitating access to large and integrated collection of individuals, characterized by a large number of variables emanating from different technologies and platforms. To achieve this goal, CEPH facilitates the setting up of integrated analyses combining clinical, genetic and environmental data, for the identification of susceptibility factors to complex multifactorial disorders Additionally, CEHP allows the reception, storage, processing and distribution of biological sample collections. At the same time, it promotes and participates in the design and setting up of genetic studies: - in partnership and full collaboration with external research groups - giving access to a large number of variables - in a sufficient number of subjects - allowing large scale integrated analyses

Synonyms: CEPH

Resource Type: institution

Keywords: environmental, genome, genetic, analysis, biological, cell, clinical, disorder, distribution, diversity, dna, human, individual, laboratory, polymorphism, process, procession, reception, research, storage, structure, subject, technology, variable

Resource Name: Centre dEtude du Polymorphisme Humain

Resource ID: SCR_008026

Alternate IDs: Wikidata: Q5464989, nif-0000-10191, ISNI: 0000 0004 0639 125X, grid.417836.f

Alternate URLs: https://ror.org/01rje3r53

Record Creation Time: 20220129T080245+0000

Record Last Update: 20260801T190343+0000

Ratings and Alerts

No rating or validation information has been found for Centre dEtude du Polymorphisme Humain.

No alerts have been found for Centre dEtude du Polymorphisme Humain.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 294 mentions in open access literature.

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

Workalemahu T, et al. (2026) Inherited genetic risk in stillbirth: A shared genomic segments analysis of high-risk pedigrees. HGG advances, 7(1), 100546.

Pennel CL, et al. (2026) Evolution of Master of Public Health Core Curriculum: Trends and Insights. Public health reports (Washington, D.C. : 1974), 141(1), 130.

Brown EA, et al. (2025) Clarification of public health pathways toward clinical careers: a pilot study describing a benchmark analysis project. Frontiers in public health, 13, 1534403.

Beneker O, et al. (2025) Urbanization and genetic homogenization in the medieval Low Countries revealed through a ten-century paleogenomic study of the city of Sint-Truiden. Genome biology, 26(1), 127.

Park JJ, et al. (2025) Key factors for successful eruption of the mandibular third molar after extraction of the mandibular second molar. Korean journal of orthodontics, 55(2), 154.

Arcos-Burgos M, et al. (2025) Neurodevelopment Genes Encoding Olduvai Domains Link Myalgic Encephalomyelitis to Neuropsychiatric Disorders. Diagnostics (Basel, Switzerland), 15(12).

Falcon A, et al. (2025) Unlocking public health competencies: the dose-response effect of problem-based learning on undergraduate student outcomes. Frontiers in public health, 13, 1560098.

Moekotte L, et al. (2025) Elevated Plasma Complement Factors in CRB1-Associated Inherited Retinal Dystrophies. Investigative ophthalmology & visual science, 66(2), 55.

Zhang R, et al. (2025) Socioenvironmental factors associated with shrimp allergy and shrimp sensitization in a large and diverse patient population and cohort study from metropolitan Detroit. The journal of allergy and clinical immunology. Global, 4(3), 100492.

Porubsky D, et al. (2025) Human de novo mutation rates from a four-generation pedigree reference. Nature, 643(8071), 427.

Perez CFM, et al. (2025) Elevated Serum Bile Acids Predict Poor Liver Outcomes in Children With Alagille Syndrome: Results From the GALA Study Group. Liver international : official journal of the International Association for the Study of the Liver, 45(12), e70423.

Gu J, et al. (2025) Identifying the genetic basis and molecular mechanisms underlying phenotypic correlation between complex human traits using a gene-based approach. bioRxiv : the preprint server for biology.

Shneider BL, et al. (2025) Plasma proteome correlations with liver stiffness in pediatric cholestasis implicate epithelial to mesenchymal transition. Hepatology communications, 9(10).

Lin A, et al. (2025) Genos: a human-centric genomic foundation model. GigaScience, 14.

Lucas-Sánchez M, et al. (2024) The Impact of Recent Demography on Functional Genetic Variation in North African Human Groups. Molecular biology and evolution, 41(1).

Plender EG, et al. (2024) Structural and genetic diversity in the secreted mucins MUC5AC and MUC5B. American journal of human genetics, 111(8), 1700.

Perez-Matas E, et al. (2024) Overexpression of BAPT and DBTNBT genes in Taxus baccata in vitro cultures to enhance the biotechnological production of paclitaxel. Plant biotechnology journal, 22(1), 233.

Kedia SK, et al. (2024) A narrative review of the CEPH-accredited bachelor's public health programs' curricula in the United States. Frontiers in public health, 12, 1436386.

Porubsky D, et al. (2024) A familial, telomere-to-telomere reference for human de novo mutation and recombination from a four-generation pedigree. bioRxiv : the preprint server for biology.

French JN, et al. (2024) Comparing the effect of imputation reference panel composition in four distinct Latin American cohorts. bioRxiv : the preprint server for biology.