Resource Summary Report

Generated by NIF on Apr 16, 2025

National Genotyping Center; Ile-de-France; France

RRID:SCR_011412
Type: Tool

Proper Citation

National Genotyping Center; Ile-de-France; France (RRID:SCR_011412)

Resource Information

URL: http://www.cng.fr/

Proper Citation: National Genotyping Center; Ile-de-France; France (RRID:SCR_011412)

Description: National genotyping center to advance research of the genetics of human diseases through internal and collaborative research programs. To this end they have developed a number of state-of-the-art genomics technology platforms and laboratories. Technology available at CNG ranges from a fully integrated high-throughput genotyping facility to a next generation sequencing platform. Activities include whole genome association studies (GWAS), pan-genomic expression profiling, epigenetic studies (DNA methylation, chromatin structure studies) and whole genome sequencing. All platforms, tools and know-how developed at CNG are open to outside groups for through collaborative research programs, and also applied by the Centre for its own research into the genetic basis of hereditary diseases.

Abbreviations: CNG

Synonyms: Centre National de Génotypage, Centre National de Genotypage, National Genotyping Center; Île-de-France; France

Resource Type: national laboratory

Funding:

Resource Name: National Genotyping Center; Ile-de-France; France

Resource ID: SCR_011412

Alternate IDs: nlx_158239

Record Creation Time: 20220129T080304+0000

Record Last Update: 20250410T070116+0000

Ratings and Alerts

No rating or validation information has been found for National Genotyping Center; Ile-de-France; France.

No alerts have been found for National Genotyping Center; Ile-de-France; France.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Soria JM, et al. (2002) A quantitative-trait locus in the human factor XII gene influences both plasma factor XII levels and susceptibility to thrombotic disease. American journal of human genetics, 70(3), 567.

Lefèvre C, et al. (2001) Mutations in CGI-58, the gene encoding a new protein of the esterase/lipase/thioesterase subfamily, in Chanarin-Dorfman syndrome. American journal of human genetics, 69(5), 1002.