

# Human population genomics and genetic epidemiology

Institut Pasteur, Paris, September 28-October 9, 2020

This two-week course presents theoretical lectures, research examples and hands-on computer training on concepts and tools used in population genomics and genetic epidemiology.

**First week - topics include:**

- Principles in genetic epidemiology and statistical genetics
- Public databases in human genomics (1,000 Genomes, ExAC, gnomAD)
- Genetic epidemiology: linkage analyses and association studies
- Genome-wide association studies (GWAS)
- Whole exome and whole genome sequencing
- Computer laboratory work in genetic epidemiology
- Research examples

**Second week - topics include:**

- Principles in population genetics
- Population structure and human demography
- Methods to detect natural selection
- Genome-wide analyses of natural selection and research examples
- Quantitative genetics and expression quantitative trait loci (eQTL) mapping
- DNA methylation
- Computer laboratory work in population genetics
- Lectures from keynote speakers

**Online registration:**

[www.pasteur.fr/en/human-population-genomics-and-genetic-epidemiology](http://www.pasteur.fr/en/human-population-genomics-and-genetic-epidemiology)

**Co-directors:**

Lluis Quintana-Murci

Aurélie Cobat

**Honorary Director:** Laurent Abel

**Practical information:**

**Deadline for application:** June 20, 2020

**Attendees:** 25 students

**Contact:** [enseignement@pasteur.fr](mailto:enseignement@pasteur.fr)

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