

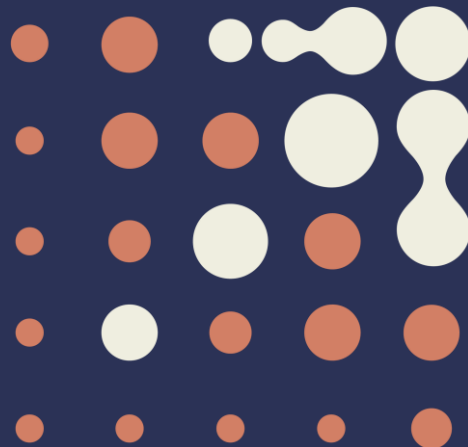
Medical Genomics course

N. Alcala & M. Foll

Rare Cancers Genomics Team

2024

International Agency
for Research on Cancer



Philosophy

Hands-on course with a focus on real-world medical problems

- > to know if you want to dedicate your professional career to medical genomics research, you need to **experience what working on a typical medical genomic project is like**
- > We will see just enough theory to grasp the main concepts and then **code and do actual data analysis**, and do **team work**

General information

Venue: IARC – WHO: the world's cancer agency <https://www.iarc.who.int>

Attendees: INSA students + CanBioS doctoral school students + IARC early career scientists (~20 students in total)

INSA | INSTITUT NATIONAL
DES SCIENCES
APPLIQUÉES
LYON



CanBioS
CANCÉROLOGIE
BIOLOGIE
SANTÉ
UNIVERSITÉ DE LYON



IARC nouveau centre in the Gerland biodistrict



Nicolas Alcala
Computational
Cancer

Genomics team,
cancer ecology
and evolution

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Matthieu Foll
Computational
Cancers

Genomics team
leader, head of
bioinformatics

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International Agency for Research on Cancer



IARC's Mission: research on cancer prevention

- ♦ .. is the **specialized cancer agency** of the World Health Organization (WHO).
- ♦ .. promotes **international collaboration** in cancer research.
- ♦ .. identifies the **causes of cancer** so that **preventive measures** may be adopted.
- ♦ .. conducts research in **LMICs** through **partnerships and collaborations** with researchers in these regions.



Main areas of IARC's research

Pillar 1 Data for action



Cancer Surveillance

Pillar 2 Understanding the causes



Nutrition & Metabolism



Genomic Epidemiology



Laboratory Support & Biobanking

Pillar 3 From understanding to prevention



Early Detection, Infections & Prevention

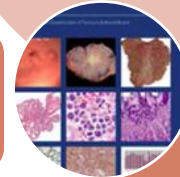


Environment & Lifestyle



Epigenomics & Mechanisms

Pillar 4 Knowledge mobilization



Evidence Synthesis & Classification



Learning & Capacity Building

The Computational Cancer Genomics team

Multi-omics characterization of tumors. Identify aggressive phenotypes.

> **Cancers of interest:** mesothelioma, neuroendocrine neoplasms (focus on lung), rare cancers (<https://rarecancersgenomics.com>)

> **Approaches:** bioinformatics, computational biology, deep-learning, modeling, evolution and ecosystems

> **Techniques:** genome, transcriptome, epigenome, single-cell and spatial omics, images



Matthieu Foll
Team leader - Computational Biol



Lynnette Fernandez-Cuesta
Team leader - Biology/Biochemistry



Nicolas Alcala
Scientist - Statistics



Laurane Mangé
PhD student - Computational Biology



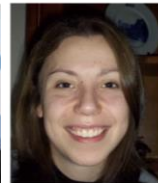
Emilie Mathian
Postdoc - Machine learning/AI



Catherine Voegelé
Data analyst - Bioinformatics



Lisa Bonheme
Postdoc - Machine learning/AI



Gabrielle Drevet
PhD student - Surgery



Lipika Kalson
PhD student

Program

- Monday November 25th
 - 11-12:00 welcome and general introduction
 - 13-15:00 genomics lecture
- Tuesday November 26th
 - 9-10:00 single-cell RNA-seq lecture
 - 10-12:00 single-cell practical part I
 - 13-14:00 spatial RNA-seq lecture
 - 14-16:00 epigenomics and multi-omic integration lecture
 - 16-18:00 single-cell practical part II
- Wednesday November 27th
 - 9-10:00 AI for digital pathology
 - 10-12:00 AI for digital pathology practical part I
 - 13-14:00 AI for multi-modal integration
 - 14-16:00 AI for digital pathology part II
 - 16-18:00 projects
- Friday November 29th
 - 9-12:00 projects
- Tuesday December 17th
 - 14-16:00 project presentations
 - 16-18:00 networking event

Practical information

Computing resources

Scientific IT platform: dedicated portal for data analysis

<http://portal.sit.iarc.fr> / HPC: `osiris.iarc.lan`

Training-MG folder on Osiris

Accounts and gateway created for **external participants**:

<https://193.51.164.141/>

login: firstname.lastname

password: SiT@Temp2024! (to be changed at first login)

Teams channel: [IARC-Medical genomics course | General | Microsoft Teams](#)

GitHub page: https://github.com/IARCBioinfo/medical_genomics_course

Lunch / coffee break

Cafeteria on ground floor

Clusters • Interactive Apps • Osiris Apps • Others Apps • My Interactive Sessions

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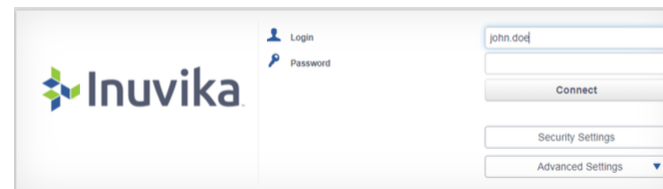
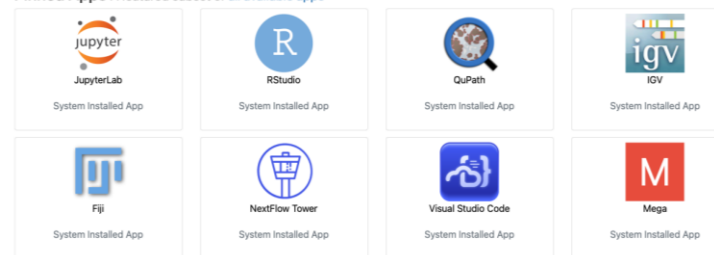
World Health
Organization



Scientific IT Portal

OnDemand provides an integrated, single access point for all of your HPC resources.

Pinned Apps A featured subset of [all available apps](#)



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Practicals

1. Single-cell and spatial transcriptomic analyses

Data on IARC's HPC Osiris

Access to jupyter or vscode through the IARC SIT platform (web browser)

2. Deep learning analysis of whole-slide pathological images

Access through Google colab notebook for access to GPU

Both using python, use links sent if needed (in particular the R to Python tutorial)

Group projects

5 projects based on actual research done at the agency

- ranging from more bioinformatics oriented to more data analysis oriented
- Groups of 3-4 students

Group presentation in December 17th (10 min + 10 min of questions per group).

Evaluation for INSA students

- grade is 50% supervisor evaluation and 50% presentation evaluation
- Criteria are: understanding of subject, clarity of explanations, organization and structure of the presentation, visuals, engagement, and answer to questions
- All group members should present something