



ALACI

LATIN AMERICAN AND CARIBBEAN ASSOCIATION OF IMMUNOLOGY

ALACI 2024 - Pre-Congress Course

3RD FRANCO-AUSTRAL COURSE ON
BIOINFORMATICS APPLIED TO
IMMUNOLOGY

COMPUTATIONAL TOOLS APPLIED TO THE ANALYSIS OF SINGLE CELL "OMICS"

31/10/2024-2/11/2024

GOAL: To provide graduate students and young researchers from Latin American countries with the fundamentals of single-cell transcriptomics and proteomics. During this course attendees will learn how to use available R-based computational and statistical methods to analyse immune cell datasets. The course is organised in such a manner that, during the mornings, the theoretical aspects of the different technologies will be discussed after listening to a keynote speaker work and during the afternoons, experts in computational and statistical methods will provide personalized training to analyse immune cell datasets (from the literature or the papers exposed during the morning). Although several virtual courses have been implemented in the last few years, we still believe that in-person meetings allow maximum interaction. Consequently, the afternoon session will be organised in a way that one professor/organiser will assist a group of 4-5 students. One assigned paper with an associated data set will be provided to each group to be analysed during the course and a final presentation of the result will be evaluated on the last day. The focus will be on implementing commonly used software and pipelines (eg CellRanger, Seurat, ArchR). We expect to foment productive, high-level discussions and to create a nurturing and supportive environment for early-career investigators and trainees that will consequently impact the quality of our research, teaching and health systems.

DIRECTED TO: The course is intended for graduate students pursuing their doctorate, as well as postdoctoral fellows and young researchers involved in research in basic, applied and clinical immunology.

Registration open until the 16th September

<https://forms.gle/MH8MDqvB8nedxnqNA>

ADMISSION PROCESS Admission will be subject to selection based on merit, geographic distribution, and gender equality among applicants. **Interested parties must send a CV and a summary of their work along with a motivation letter that justifies the need to implement these tools in their research work. Candidates must demonstrate knowledge and experience in using the "R" programming language and a good understanding of English**

MINIMUM AND MAXIMUM NUMBER OF STUDENTS:

Minimum: 20

Maximum: 40

COST:

Members of a national immunology society belonging to ALACI (<https://www.alaci.org/en/sociedades-afiliadas/>), AINCA members, or participants from low or middle-income countries according to the World Bank Classification: 40 US dollars.

Non-ALACI members, non-AINCA members, or persons from high-income countries according to the World Bank Classification: 60 US dollars.

FELLOWSHIPS AND TRAVEL GRANTS

1. ALACI MEMBERS assisting ALACI 2024 will not pay the registration fee.
2. There are 12 accommodation fellowships available for ALACI MEMBERS assisting this course considering academic and scientific training, geographical distribution, and gender balance. Please, indicate if you need accommodation in the registration form.
3. There are 10 travel grants (500 euros; Bs As-Santiago or vice versa) that will be accessible exclusively to master students, PhD students or postdoctoral researchers from South American universities and research institutions, wishing to participate in these two courses: the 3rd Franco-Austral Course on Bioinformatics preceding the ALACI conference (Buenos Aires, Argentina) and the 7th Advanced Course on Immunity and Cancer Immunotherapy (Santiago, Chile). The award will be reimbursed in Chile, after finishing both events. For more information and to access to this travel grants, please register in the following form:
<https://forms.gle/N4DMNDbF8MVGzRSH8>.
4. Postulation to both benefits (the accommodation and the travel grants) are accepted

SCIENTIFIC ORGANIZERS

Dr. Eliane Piaggio

Translational Immunotherapy
U932.Immunity and Cancer
Institut Curie
26 rue d'Ulm
75248 Paris cedex 05 - France
Tel.+33 (0)1 56 24 55 00
e-mail: eliane.piaggio@curie.fr

Dr. Mariana Maccioni. FCQ-UNC.

Departamento de Bioquímica Clínica
Facultad de Ciencias Químicas. CIBICI-CONICET.
Universidad Nacional de Córdoba.
Haya de la Torre y Medina Allende. Ciudad
Universitaria. Córdoba. Argentina
e-mail: mariana.maccioni@unc.edu.ar

Dr Joshua Waterfall

Integrative Functional Genomics of Cancer
U830. Cancer,Heterogeneity, Instability and
Plasticity
Institut Curie
26 rue d'Ulm
75248 Paris cedex 05 - France
Tel.+33 (0)1 56 24 55 00
e-mail: joshua.Waterfall@curie.fr

Dr. Alvaro LLadser

Principal Investigator
Fundacion Ciencia & Vida - Universidad San
Sebastián
Avda. del Valle Norte N° 725
Huechuraba, Santiago, Chile
e-mail: alladser@cienciavida.org

Coordinators

Dras. Carolina Jancic-Mercedes Borge

Academia Nacional de Medicina. Jefe de Trabajos Prácticos. Departamento de Microbiología,
Parasitología e Inmunología. Facultad de Medicina. UBA. Imex- Conicet

Teaching Staff

Dr. Nicolás Gonzalo Nuñez

Departamento de Bioquímica Clínica.
CIBICI-CONICET
Facultad de Ciencias Químicas. UNC

Dr. Jimena Tosello

Translational Immunotherapy
U932.Immunity and Cancer
Institut Curie, París

Dr. Dario Rocha

Translational Immunotherapy
U932.Immunity and Cancer
Institut Curie, París

Dr. Andres Hernandez Oliveras

Fundación Ciencia & Vida
Facultad de Medicina y Ciencia, Universidad
San Sebastián

Joaquín Merlo

Laboratorio de Glicómica Funcional y Molecular
(Programa de Glicociencias),
IBYME – CONICET

Yamil Mahmoud

Laboratorio de Glicómica Funcional y Molecular
(Programa de Glicociencias)
IBYME – CONICET

Dr. Tomás Dalotto

Laboratorio de Glicomedicina
IBYME – CONICET
Buenos Aires

TENTATIVE PROGRAM

Day 1	
09:00 a 9:45	General introduction and presentation of the course. Presentation of the participants
09:45 a 10:15	Overview of gene regulation-Applications of scRNA/TCRseq/ATAQ seq
10:15 a 10:45	General concepts on how to run a scRNA/TCR/ATAQ seq technologies
10:45 a 11:00	Coffee Break
11:00 a 13:00	Theoretical concepts on how to run a scRNA
13:00 a 14:30	Lunchtime
14:30 a 16:30	Bioinformatics analysis of single cell transcriptomics
16:30 a 18:30	“Hands on computer training”
Day 2	
09:00 a 10:45	sc technologies: Introductory concepts. ATACseq-multiomics-spatial transcriptomics Biological and technological introduction QC and filtering Dimensionality reduction and visualization Clustering and peak calling Gene score prediction and motif analysis
10:45 a 11:15	Coffee Break
11:15 a 12:00	General concepts of T cell biology and TCR-

12:00 a 13:00	Hands on training TCR data analysis
13:00 a 14:30	Lunchtime
14:30 a 15:30	Hands on training TCR data analysis-
15:30 a 16:00	Coffee Break
16:00 a 18:30	Hands on training TCR data analysis
Day 3	
9:00 a 10:30	Spectral flow cytometry. Basic concepts of full spectrum cytometry. Deconvolution. Demonstration of autofluorescence extraction. Spectral data analysis applying dimensionality reduction. CITE- Seq.
10:30 a 11:00	Coffee Break
11:00 a 13:00	“Hands on computer training”
13:00 a 14:30	Final Conclusions