









THE 2018 EXPOSOME SHORT COURSES SERIES

EXPOSOME PRACTICE

ANGLET 25-29 JUNE, 2018





EXPOSOME BASIS: UTRECHT JULY



SURF64 SUMMER SCHOOL:

STATISTICS AND MACHINE LEARNING FOR OMICS PROFILING AND INTEGRATION IN EXPOSOME RESEARCH













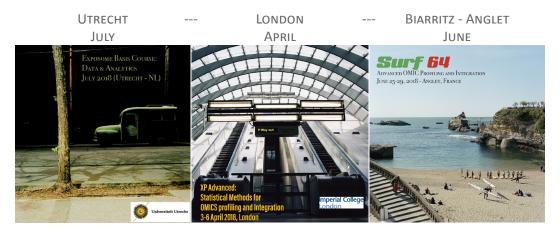








THE 2018 EXPOSOME SHORT COURSES SERIES



THE EXPOSOME SHORT COURSES SERIES is a multi-institution programme on Statistical models and Bioinformatics tools to analyse Exposome data. It comprises three complementary one-week courses:

- EXPOSOME BASIS (UTRECHT UNIVERSITY, NL): the Molecular Epidemiology and Exposome Course (MEEC) is an introductory course providing a comprehensive introduction to the concept of the exposome and its practical implementation. The course focuses on OMICs data, their features, and the challenges their statistical analysis raises. The MEEC proposes a series of lectures, seminars and practical describing the main statistical methods used in molecular epidemiology. These include univariate models and multiple testing correction strategies (FWER, FDR), dimension reduction techniques, and variable selection approaches.
- EXPOSOME ADVANCED (IMPERIAL COLLEGE, UK): XP ADVANCED is an advanced course presenting further techniques to analyse and integrate OMICs data in an exposome concept. The course build upon a statistical background such as the one taught in the MEEC course to introduce necessary extensions of these methods in order to (i) accommodate complex study designs; (ii) Improve results interpretability; and (iii) handle multiple sets of OMICs data. In addition to regression-based profiling approaches, XP ADVANCED also features the exploration of machine learning techniques, including network inference and their practical application to OMICs data. The course will develop the theoretical background of these methods and their applicability to OMICs data in Exposome Sciences.
- EXPOSOME PRACTICE (IMPERIAL COLLEGE UNIVERSTIY OF PAU ANGLET): SURF 64 is a
 one-week summer school focusing the on the practical application of all methods and
 principles developed in the MEEC and XP ADVANCED to real data. The summer school
 will consist in 5-days supervised group work using real data sets and addressing real
 research questions on OMICs analysis, interpretation and integration. The course also
 includes a series of lectures, seminars, and tutorials illustrating solutions to OMICs
 profiling and integration in a real-life setting.



















OVERVIEW

SURF 64 will take place at the UPPA-Anglet Campus in the Basque country and will run from the 25th to the 29th June, 2018. The course comprises a 5 full days of group-work on real data analysis (including multi omics data) addressing real research questions. These projects will include multi-OMICs data measured in the same individuals and their analysis will call upon multivariate techniques as well as the use of network models and companion bioinformatics tools as taught in the MEEC and XP-Advanced courses. Datasets will be provided by course organisers but participants can also submit a dataset they would like to see analysed during the course. Project based work will be complemented by seminars/lectures related to the methods used in the group-work.

PROJECT WORK

At the beginning of the course a list of research projects will be proposed. Each of these projects will (i) address an original research question, and (ii) include multi-omic data. Attendees will form groups and will chose a project they will work on throughout the course. This project-based work will be supervised by the project leaders on a daily basis, and technical guidance will be provided both by course facilitators and lecturers for specific methodology. The last afternoon of the course will be dedicated to presentations (seminar format) where each group will present their research questions and their main results to their fellow participants and a panel of experts.

LEARNING OUTCOMES

After SURF 64, students will have practical experience in:

- Multi-OMIC profiling using univariate, multivariate approaches and their extensions
- implementing these approaches to analyse real-life data
- integrating different OMICs data and interpreting results using established bioinformatics tools
- inferring network topologies for results interpretation and feature selection
- providing results in a reproducible and sustainable manner using open source tools; and will have been introduced to machine learning/deep learning.

















TARGET AUDIENCE

SURF 64 will be of interest to academics (students, and researchers), and scientists from the industry (pharmaceutical companies, insurance companies, food industries...). Experience in statistics, OMICs data and use of R statistical software is required (e.g. MEC-StatXP courses, Advanced XP course).

Participants should bring their own laptops, and could submit a dataset they would like to analyse during the school.

Up to 30 participants can register.

REGISTRATION/FEES

REGISTRATION can be done online:

<u>http://www.imperial.ac.uk/school-public-health/study/short-courses/XP-series/</u>
For any question please send an email to:

m.chadeau@imperial.ac.uk or benoit.liquet@univ-pau.fr

Course fees:

Early bird registration (until April 30th 2018)
 Academic: £1,000 - Non-academic: £1,200

• Standard registration

Academic: £1,100 - Non-academic: £1,400

LOCATION

VENUE: UPPA ANGLET CAMPUS – PARC MONTAURY

Teaching will take place at the Biarritz-Anglet Campus of UPPA.



ACCOMMODATION

ACCOMMODATION: STUDENTS RESIDENCE EUGÈNE GOYHENECHE

Accommodation is not included in the course fees, but negotiated rental fares are available for flats in Bayonne Student Residence: £200 for the rental of a studio flat (limited availability).

Flats feature one small kitchenette and a shower room, and are located at a walking distance from Bayonne city centre and less than 1.5km from the campus.

Interested attendees should contact the organisers to check availability and books their flat.

MRC-PHE
Centre for Environment & Health

















CONTRIBUTORS

IMPERIAL COLLEGE LONDON (UK):

DR MARC CHADEAU-HYAM, Senior Lecturer in Statistical Bioinformatics, Dept of Epidemiology and Biostatistics (EBS). Honorary Reader, Utrecht University.

Dr Sabrina Rodrigues, Research Associate, EBS

DR MARYAM KARIMI, Research Associate, EBS.

MRS BARBARA BODINIER, Research Assistant, EBS.

University of Pau et Pays de l'Adour, (FR):

PROF BENOÎT LIQUET, Professor in Statistics, Member of LMAP. Affiliated to ACEMS, Queensland University Technology.

UTRECHT UNIVERSITY (NL):

PROF ROEL VERMEULEN, Professor, Institute for Risk Assessment Sciences (IRAS). Honorary Professor, Imperial College London.

DR LÜTZEN PORTENGEN, Senior Scientist, IRAS.

DR JELLE VLAANDEREN, Assistant Professor, IRAS.

EMORY UNIVERSITY (US):

PROF GARY MILLER, Rollins School of Public Health

INSTITUT NATIONAL DE LA RECHERCHE AGRONOMIQUE (INRA-AGROPARISTECH) (FR):

DR JULIEN CHIQUET, Senior Researcher, Dept of Statistics

INSTITUT NATIONAL POUR LA RECHERCHE MEDICALE (INSERM) (FR):

DR CYRILLE DELPIERRE, Senior Researcher, Université de Toulouse

INSTITUT PASTEUR (FR):

DR BENNO SCHWIKOWSKI, Group Head, Systems Biology Group

















Monday 25,

REFRESHER ON METHODS AND PROJECTS DEFINITION:

JUNE 9.15-09:30 WELCOME AND REGISTRATION

9.30-10:30 LECTURE: Refresher on OMICs Profiling and Integration

Speaker: M Chadeau-Hyam

10:30-11:00 Datasets overview – projects allocation

Speaker: J Vlaanderen

11:30-13:00 Tutorial: Reproducible results using RStudio-RMarkdown and GitHub

Speaker: B Liquet

14:00-18:00 GROUP WORK: Data exploration – Repository set up – Analytical Plan

TUESDAY 26,

JUNE EXPLORATORY ANALYSES - VISUALISATION

9:00-10:00 GROUP PRESENTATION: Projects overview and Analytical plan

10:00-11:00 Tutorial: Big Data Visualisation and Exploration

Speaker: G Miller

11:00-13:00 Group Work: Exploratory analyses – Association Studies

14:00 -15:00 SEMINAR: Resampling techniques: Calibration/Validation/Stability

Speaker: L Portengen

15:00-18:00 GROUP WORK: Visualise and interpret profiling results

WEDNESDAY 27,

JUNE SENSITIVITY/STABILITY ANALYSES – RESULTS INTERPRETATION

09:30-10:30 Group Presentation: Preliminary results – planned refined analyses

10:30-12:00 Tutorial: Pathway Exploration – Bioinformatics tools

Speaker: G Miller

13:00-14:00 SEMINAR: Introduction to Cytoscape and Sub-Network Models

Speaker: B Schwikowski

14:15-17:30 GROUP WORK: Implementation of stability analyses, Results generation

and interpretation

















THURSDAY 28,

OMICS INTEGRATION:

JUNE 09:30-10:30 GROUP PRESENTATION: Finalised results & plans for OMICs integration

10:45-12:15 LECTURE: Introduction to Network Inference in practice

Speaker: J Chiquet

12:15-13:00 SEMINAR: The role of advanced statistics in Exposome Sciences

Speaker: R Vermeulen

14:00-15:00 Lunch Seminar: Variational inference in the Poisson-lognormal model:

applications to multivariate analysis of count data

Speaker: J Chiquet

15:00-18:00 GROUP WORK: Implementing network models for (multi)-OMIC data

FRIDAY 29,

JUNE NETWORK MODELS IN PRACTICE

09:30-10:30 LECTURE: Epidemiological approaches to results interpretation:

confounding, mediation, temporality and causation

Speaker: C Delpierre

10:30-13:30 Group work: finalising analyses & presentation 13:45 -14:30 LUNCH SEMINAR: Mechanistic investigation

Speaker: M Chadeau-Hyam

14:30-17:00 FINAL GROUP PRESENTATION















