

# An Introduction to Omics

## Two-day training course

A two-day course aimed at familiarizing participants with the basis and application of various omics disciplines: genomics, transcriptomics, metabolomics, proteomics, and bioinformatics. Each of the omics disciplines will be covered by a lecture and a practical bioinformatics session. By the end of the course users should understand, for each omics level: the basis of the discipline, the instrumentation used to generate high-throughput biological data, key applications, and how to visualise the resulting data using commonly used software packages. Participants will also be aware of how different large-scale data sets can be integrated in order to obtain better biological inference, and appreciate the nature of other modern challenges in bioinformatics.

### Dates

3–4 March 2014

### Time

DAY 1: 10:00am–5:00pm

DAY 2: 9:00am–5:00pm

### Course fee

£300

### Target group

Research students and staff who wish to deepen their understanding of high-throughput data generation and analysis.

### Venue

Jura teaching lab, Level 4 Annexe,  
University Library

### Speakers

Mike Barrett, Richard Burchmore,  
Karl Burgess, Graham Hamilton,  
Pawel Herzyk

### Registration and enquiries

Contact: Tanita Casci

[Tanita.Casci@glasgow.ac.uk](mailto:Tanita.Casci@glasgow.ac.uk)

### Further information

Online at: [www.polyomics.gla.ac.uk/  
services.html#gpTraining](http://www.polyomics.gla.ac.uk/services.html#gpTraining)

Follow us on Twitter: [@polyomics](https://twitter.com/polyomics)

Glasgow Polyomics runs formal and informal training session for internal and external users. Informal training, including one-on-one tutorials, can be arranged on request by contacting Tanita Casci.

### Programme Day 1

#### Jura teaching lab, Level 4 Annexe, University Library

##### Overview of Polyomics (Mike Barrett)

10:00am–10:30am

##### Tea/Coffee break

10:30am–10:45am

##### Genomics (Graham Hamilton)

10:45am–11:45am: Lecture

11:45am–12:45pm: Practical session

##### Lunch

12:45pm–1:45pm

##### Transcriptomics (Pawel Herzyk)

1:45pm–2:45pm: Lecture

2:45pm–3:45pm: Practical session (Part 1)

##### Tea/Coffee break

3:45pm–4:10pm

##### Transcriptomics (cont'd)

4:10pm–4:40pm: Practical session (Part 2)

##### Questions and discussion: Genomics & Transcriptomics (Pawel Herzyk)

4:40pm–5:00pm

### Programme Day 2

#### LC-MS- and NMR-based Metabolomics (Karl Burgess)

9:00am–10:30am: Lecture

10:30am–11:30am: Practical session

##### Tea/Coffee break

11:30am–11:45am

##### Proteomics (Richard Burchmore)

11:45am–12:45pm: Lecture

##### Lunch

12:45pm–1:45pm

##### Proteomics (cont'd)

1:45pm–2:45pm: Practical session

##### Integrating multiple datasets: a case study (Mike Barrett)

2:45pm–3:45pm

##### Tea/Coffee break

3:45pm–4:00pm

##### Biomarker discovery (Mike Barrett)

4:00pm–4:40pm

##### Closing remarks and Discussion (Mike Barrett)

4:40pm–5:00pm

