

# MS203-15 Patterns and Populations

**23/24**

**Department**

Warwick Medical School

**Level**

Undergraduate Level 2

**Module leader**

Sascha Ott

**Credit value**

15

**Module duration**

10 weeks

**Assessment**

100% coursework

**Study locations**

University of Warwick main campus, Coventry Primary

field trip location: FSC Dale Fort ( <https://www.field-studies-council.org/locations/dalefort/> )

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## Description

### Introductory description

This module focuses on how complex biological systems can be described and analysed using mathematical and computational approaches. For example, (1) new sequencing data is revealing insight into the connectivity in gene regulation; and (2) mathematical models underly key concepts in population and community ecology. Such information is critical as we look to understand the effects of the Anthropocene on species' genetic diversity and extinction.

The module will be compulsory for second year Integrated Natural Sciences (INS) students. It is a continuation of MSci Integrated Natural Sciences Year 1 laboratory modules that focus on molecules, cells and organisms to further increase scale to population level. The programme extends the first year of the degree using more advanced methods and analysis, but retaining the Integrated Natural Sciences ethos, solving problems by drawing freely from the methods and mindsets of more than one discipline.

### Module aims

The module will focus primarily on computational methods to analyse data, model interactions and spatial patterns and link changes at the DNA and transcriptional level to outcomes in spatial developmental patterns, species interaction and population ecology. The students will be equipped to extract quantitative data from raw sequencing reads, derive network information and

mathematical models to describe multidimensional patterns and complex multi-species interactions.

## **Outline syllabus**

This is an indicative module outline only to give an indication of the sort of topics that may be covered. Actual sessions held may differ.

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Block 1 (weeks 1 - 4): Genes and gene network analysis of high-throughput sequencing data. How to handle data, detect and correct biases, and assure data quality. How to extract differential expression and gene network information.

Block 2 (weeks 5 - 7): Ecology and evolution in populations, communities, and ecosystems. Using the sequencing and data analysis tools developed in Block 1 to characterise real-world communities. Investigate feedback(s) across scales, how species interactions scale-up to influence ecosystem properties and vice versa how ecosystems drive or constrain the evolution of species

The module ends with a field trip to collect data and perform statistical analysis.

## **Learning outcomes**

By the end of the module, students should be able to:

- Utilise software packages for analysing sequencing data (Block 1)
- Comprehend concepts underlying transcriptomics data and their representation (Block 1)
- interpret and apply data to understand network interactions in ecology and evolution of biological systems (Blocks 1-2)
- Apply statistical knowledge to estimate uncertainties in analysis (Blocks 1-2)
- Apply concepts such as species interactions and feedback to ecological systems (Block 2)
- Present concisely, appropriately and effectively on topics discussed in class (Blocks 1-2)
- Synthesise biological concepts related to network interaction (Blocks 1-2)
- Understand how to use the literature to deepen understanding (Block 1-2)
- Evaluate biological data using different techniques and able to critique the appropriateness of each methodology

## **Indicative reading list**

Michael Begon & Colin R. Townsend. Ecology: From Individuals to Ecosystems. 5th Ed.

## **Interdisciplinary**

The students will apply computational and mathematical methods to understand and describe complex biological phenomena.

## **Subject specific skills**

Knowledge of key methods in bioinformatics, data handling and analysis. Knowledge how to apply analysis tools to extract quantitative data from nucleotide reads of biological samples. Knowledge about how to interpret data in the context of ecological and evolutionary theory.

## Transferable skills

Students will be able to demonstrate integrated thinking across the Sciences. Oral presentations on quantitative methods applied to biological problems.

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## Study

### Teaching split

Provider	Weighting
Life Sciences	50%
Warwick Medical School	50%

### Study time

Type	Required
Lectures	8 sessions of 1 hour (5%)
Practical classes	7 sessions of 2 hours (9%)
Fieldwork	5 sessions of 8 hours (27%)
Private study	58 hours (39%)
Assessment	30 hours (20%)
Total	150 hours

### Private study description

Reading around topics and learning tools to perform analyses. Also includes deepening understanding of statistical approaches.

### Costs

No further costs have been identified for this module.

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### Assessment

You do not need to pass all assessment components to pass the module.

### Assessment group A

	<b>Weighting</b>	<b>Study time</b>
Block 1 Presentation Presentation related to bioinformatic analysis	25%	7 hours
Bioinformatic data analysis Students complete data analysis, evaluation and interpretation of data, based on techniques taught in the block.	25%	8 hours
Block 2 presentation Oral presentation in Block 2 on a topic set by block lead	25%	7 hours
Field trip report Students complete data analysis, evaluation and interpretation of data, based on the field trip.	25%	8 hours

### **Feedback on assessment**

Oral feedback on presentations and for first report.

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## **Availability**

### **Courses**

This module is Core for:

- Year 2 of UMDA-CF10 Undergraduate Integrated Natural Sciences (MSci)