

Thursday 29 – Friday 30 July 2021

Workshop overview

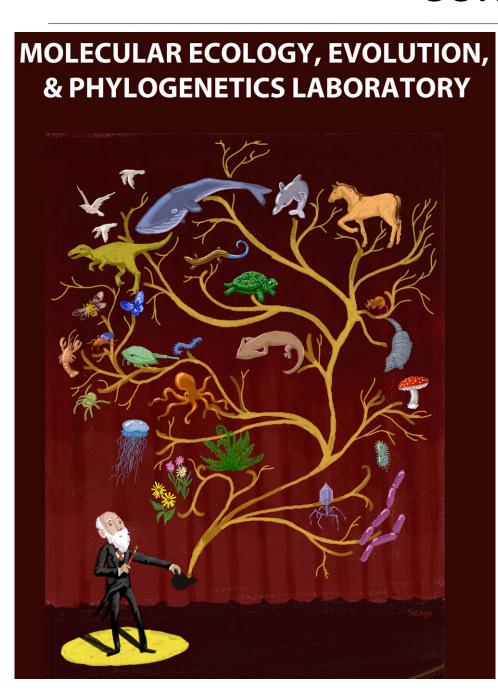
- Interpreting phylogenetic trees
- Phylogenetic methods
- Bayesian phylogenetic analysis
- Molecular dating
- Phylogenomics







Contributors



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Programme – Day 1

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11.00 – 11.10 Welcome

11.10 – 11.50 Lecture 1.1: Introduction to molecular phylogenetics

12.00 – 12.45 Lecture 1.2: Evolutionary models

13.00 – 13.45 Lecture 1.3: Phylogenetic data

14.00 – 14.45 Lecture 1.4: Phylogenetic methods

15.00 – 15.45 Seminar: Phylogenetic analysis with IQ-TREE

15.45 – 16.00 Additional time for questions
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Programme – Day 2

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11.00 – 12.00 Lecture 2.1: Bayesian phylogenetic analysis
12.15 – 12.45 Lecture 2.2: Models and priors
13.00 – 13.45 Lecture 2.3: Molecular dating
14.00 – 14.30 Lecture 2.4: Calibrating the molecular clock
14.45 – 15.15 Lecture 2.5: Phylogenomics
15.15 – 16.00 Additional time for questions
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Lecture materials

- Lecture slides available on Github github.com/simon-ho/SydneyPhyloWorkshop
- Recordings of talks will be available on Dropbox after the workshop

Practical exercises

Work through practical exercises in your own time

- Install software on your own computer
- Data files available on Github
- Brief video introduction provided (Dropbox)
- Answers to practical exercises provided (Dropbox)

- **Practical 1.1** Sequence alignment and phylogenetic analysis using *MEGA*
- **Practical 1.2** *IQ-TREE* workshop tutorial
- **Practical 2.1** Bayesian analysis and molecular dating using *BEAST*
- **Practical 2.2** Phylogenomic analysis using *ASTRAL*

Zoom webinar format

- Submit questions during webinar
 - Q&A function (questions not visible to attendees)
 - Chat function

- Answer to questions:
 - Typed responses during or after talks
 - Verbal responses after talks