

Thursday 29 – Friday 30 July 2021

# Workshop overview

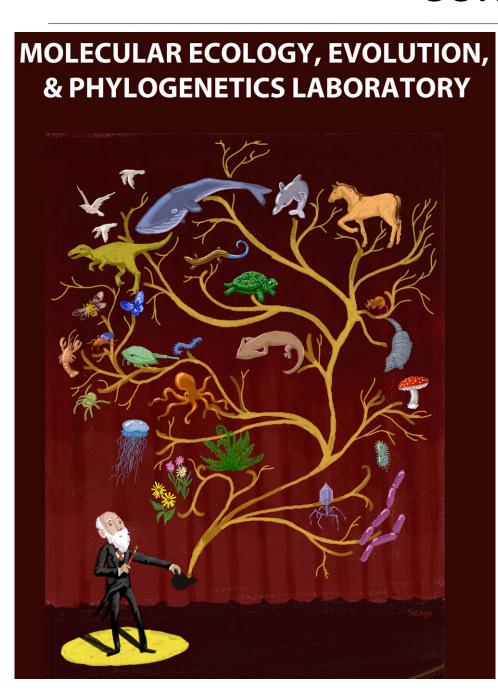
- Interpreting phylogenetic trees
- Phylogenetic methods
- Bayesian phylogenetic analysis
- Molecular dating
- Phylogenomics







#### Contributors



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## Programme – Day 1

```
Welcome
11.00 - 11.10
11.10 - 11.50
               Lecture 1.1: Introduction to molecular phylogenetics
               - Break -
12.00 – 12.45 Lecture 1.2: Evolutionary models
13.00 - 13.45
              Lecture 1.3: Phylogenetic data
               - Break -
14.00 - 14.45
               Lecture 1.4: Phylogenetic methods
               Seminar: Phylogenetic analysis with IQ-TREE
15.00 - 15.45
15.45 – 16.00 Additional time for questions
```

## Programme – Day 2

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11.00 - 12.00
              Lecture 2.1: Bayesian phylogenetic analysis
               - Break -
12.15 – 12.45 Lecture 2.2: Models and priors
13.00 – 13.45 Lecture 2.3: Molecular dating
               - Break -
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
```

#### 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