Evolutionary Thinking 2022 TA session week 1 – phylogeny tree

Jilong Ma aujilongm@birc.au.dk





Outline

1. Short Intro (15 minutes)

Myself

Grouping!

2. Learning outcome of this week (20 minutes)

Tree/Phylogeny: Interpretation

Sequence to Phylogeny

3. Working on Tree Pretest (30 minutes)
Going through solutions (15 minutes)

4. Install MEGA 11 Explore a bit ©





About Myself

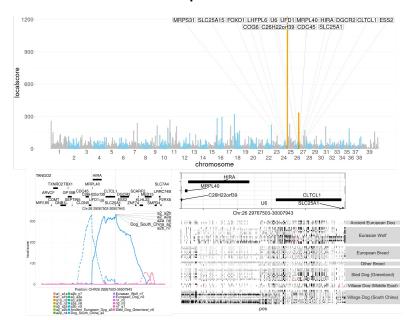
Jilong Ma (Chi lung)

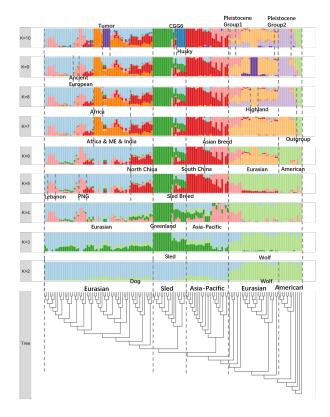
MSc in Bioinformatics, University of Copenhagen 18-20

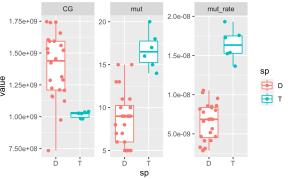
Second year PhD student with Mikkel H Schierup.

My work experience with

Population Genetics
Dogs and wolves
Comparative Genomics
Social spiders









A unique model for social transition

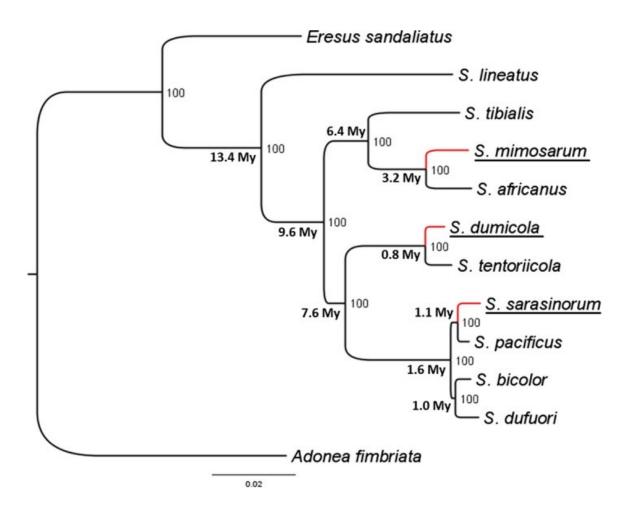


A social spider species. Stegodyphus dumicola





A unique model for social transition



Stettepani et.al 2016





Assign Groups

We have 30 registered students
Assign 8 study groups (3-4 people each)
---General TA session exercises/paper discussion
---Hand-in report as a group





Assign Groups

We have 30 registered students
Assign 8 study groups (3-4 people each)
---General TA session exercises/paper discussion
---Hand-in report as a group

Arrange yourself by birthday, like

Jan 1st, Feb 3rd, Feb 19th, ..., Dec 31st

Let's hope there is no strong correlation between birthday and study programme

Go to Brightspace - Course Tools - Group for signing in the group





Learning outcome of this week

Reading phylogeny trees (Wednesday)

Terminology for phylogeny

What is it describing?

Common mis-understanding?

Building a phylogeny tree step by step (Friday)

From sequence to phylogeny

Sequence alignments (pairwise, multiple sequence alignment)

Distance matrix and substitution models.

Phylogeny building algorithms





Learning outcome of this week

Reading phylogeny trees (Wednesday)

Terminology for phylogeny

What is it describing?

Common mis-understanding?

Building a phylogeny tree step by step (Friday)

From sequence to phylogeny

Sequence alignments (pairwise, multiple sequence alignment)

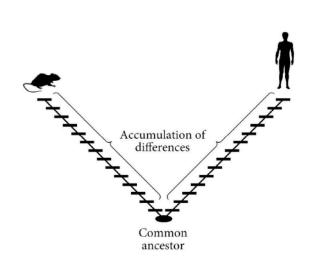
Distance matrix and substitution models.

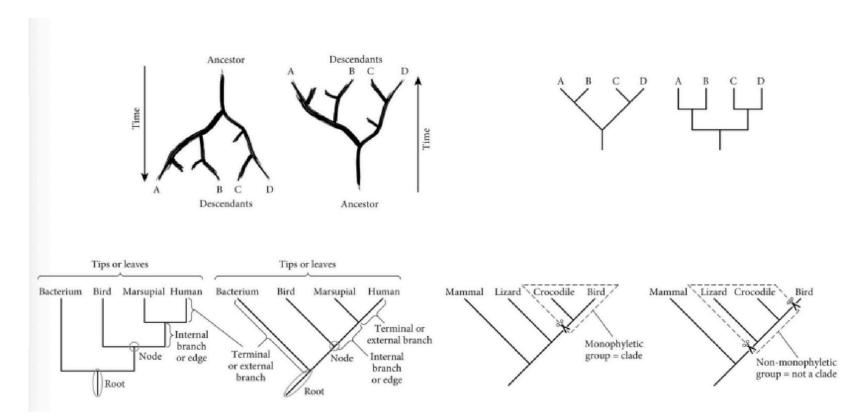
Phylogeny building algorithms





Terminology - Recap









Discussion

Reading phylogeny trees (Wednesday)
 What is it describing?
 Common mis-understanding/pitfalls?





