

# Applications: Sequencing

**BIOL** 432

#### Group assignments





Group 1: puRple loosesRife: #93748a

Hayley, Vivian, Sehaj, Anish, Minji

Group 2: GitGood: #ffooff

Kai, Pallavi, Liam, Sarah, Nikita

Group 3: Lustrous Loosestrifes: #9900cc

Claire, Jamie, Nestor, Zoe, Jonas

#### Research Background

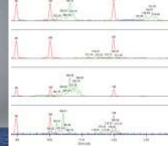


#### University of Windsor & GLIER – Bsc (Hs), MSc











**Hugh MacIsaac** 

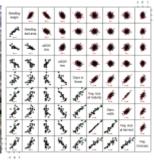
Dan Heath

#### University Toronto – PhD











Spencer Barrett

#### Research Background



Duke University – Postdoc







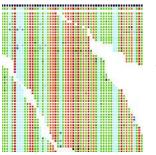




Tom Mitchell-Olds University of British Columbia – Postdoc





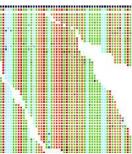


Vancouver, BC

Loren Rieseberg
University of Tuebingen – Postdoc











Oliver Bossdorf





# Rapid Evolution

in novel environments



## Ecology & Evolution in the Anthropocene

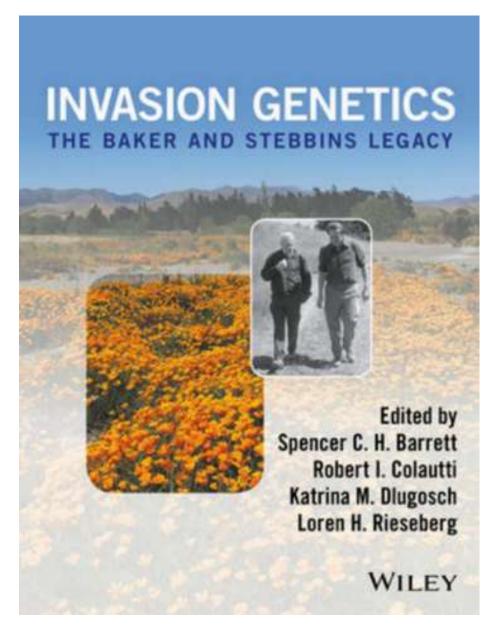




Environment --> Natural Selection --> Genome Evolution

#### **Invasion Genetics**





"Invasion genetics of the spiny waterflea"

- Colautti et al. 2005

"Invasion genetics is a relatively new discipline that investigates patterns of genetic variation in populations of invasive species and their ecological and evolutionary consequences."

- SCH Barrett 2016

#### Ecological and environmental genomics



"Invasion genetics is a relatively new discipline that investigates patterns of genetic variation in populations of invasive species and their ecological and evolutionary consequences."

- SCH Barrett 2016

**Ecological & environmental genomics** investigate patterns of genome-wide variation in natural populations or species communities, to address ecological and environmental questions.



## Methods

## Field experiments





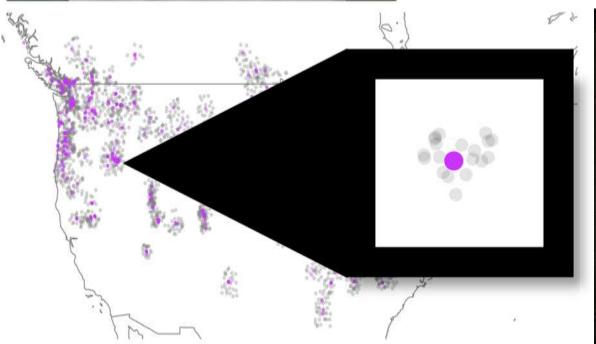
## Computation, bioinformatics and data science







- > install.packages("baRcodeR")
- > library(baRcodeR)





#### eDNA & DNA barcodes for environmental monitoring



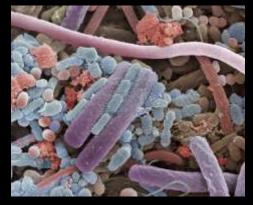




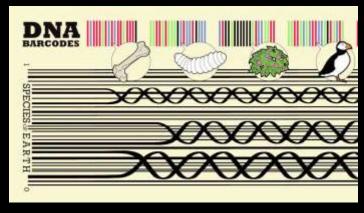








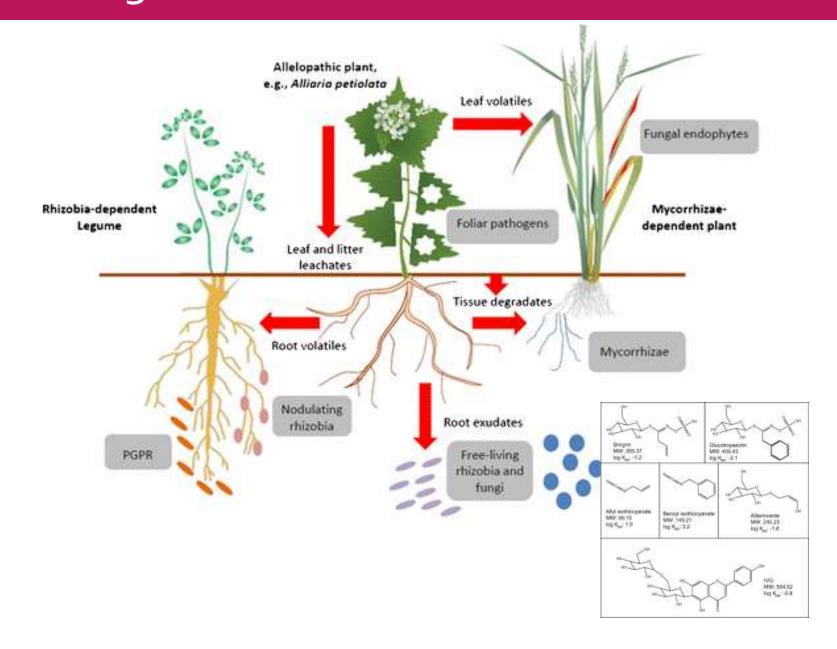
Barcode of Life Project www.boldsystems.org



https://www.youtube.com/watch?v=ZImiXgU6bCk

## Metagenomics of the soil microbiome



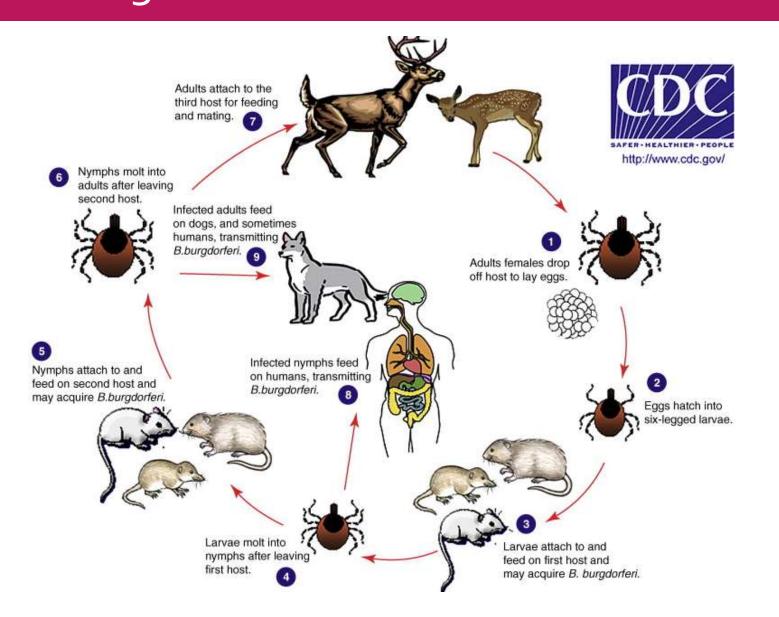




#### Metagenomics of ticks & their microbiomes









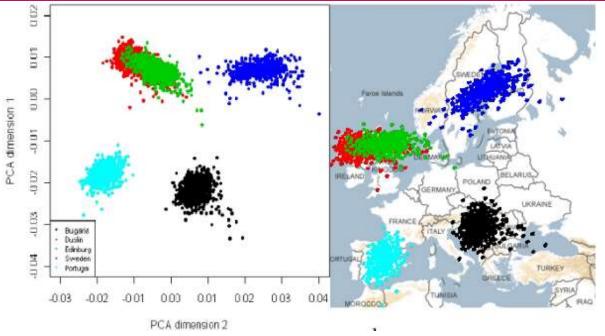




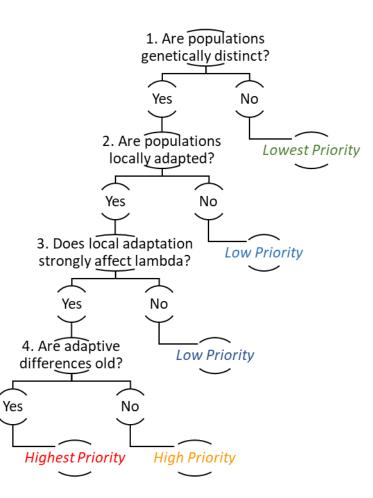


## Population genomics









#### Teaching approach – 3 pillars



#### 1. Learn by doing

Hands-on tutorials

Independent assignments

Group project

#### 2. Emphasis on transferrable skills

Coding

Data Science (collect  $\rightarrow$  manage  $\rightarrow$  visualize  $\rightarrow$  analyze  $\rightarrow$  report)

Communication

Teamwork

3. Cumulative learning – each day builds on previous activities/tutorials

## QUIZ – What are the three pillars of our approach?



1.

2.

### **Group Introductions**



Name

Program

Future Goals (short-term and long-term)

Group team name and hexadecimal colour code

Group member who speaks the most languages