1st EvoGenomics Methods Workshop

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Course intro: logistics

• GitHub repository with everything:

https://github.com/eachambers/EvoGeno-Methods-Workshop

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- GitHub repository with everything:
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- Schedule: Fridays from 10AM-1PM

Date	Topic	Instructor
April 21	Basics and QC	Anne
April 28	Basics of selection and demographic analyses	Phred
May 5	Comparative genomics	Yocelyn

Course intro: logistics

- Structure of each workshop:
 - Lectures
 - Exercises
 - Go through exercise together
 - All on GitHub! We'll post answers afterwards

Topics

- 1. Basics of genomics (slides here)
 - · Quick summary of approaches: library prep & bioinformatics assembly (standard filtering, mapping, and variant calling)
 - · How do methods compare? How does it depend on the questions you want to ask?
 - Considerations and biases for different genomic data types
 - · Brief review of sequencing/input/output file types
 - Exercises here

Course intro: other stuff

- This is the first time we're doing this
- If this stuff is all new to you:
 - Good judgement comes from experience, experience comes from bad judgement, so make mistakes!
 - Take breaks to avoid frustration
 - Keep a "lab" notebook
 - Save a copy of unedited, raw data
- If this stuff is all familiar to you:
 - Please help your peers! Use this as teaching experience
- Be respectful and supportive!