

RNA-seq data analysis workshop

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06/14/21



Who are we? The Data Analytics Core



Mission statement

Facilitate advanced genomic & bioinformatic data analysis solutions to CQB faculty & the Dartmouth research community



James O'Malley, PhD
Director



Shannon Soucy, PhD Senior Research Scientist



Tim Sullivan, BABioinformatics Research Scientist



Owen Wilkins, PhD
Bioinformatics Research Scientist

- > Genomic data analysis
 - Analytical support for Dartmouth researchers
 - Pipeline development & maintenance
- > Bioinformatics consulting
- > Publication & grant support
 - Writing for methods & results sections
 - Letters of support
- Training
 - Group workshops

Workshop outline



Millions/billions of reads from sequencing

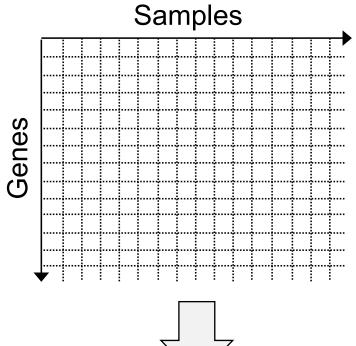


FASTQ file:

@SRR1039508.1 HWI-ST177:290:C0TECACXX:1:1101:1225:2130 length=63 CATTGCTGATACCAANNNNNNNGCATTCCTCAAGGTCTTCCTCCTTACGGAATTACA

HJJJJJJJJJJJJJJHHHFFFFFD

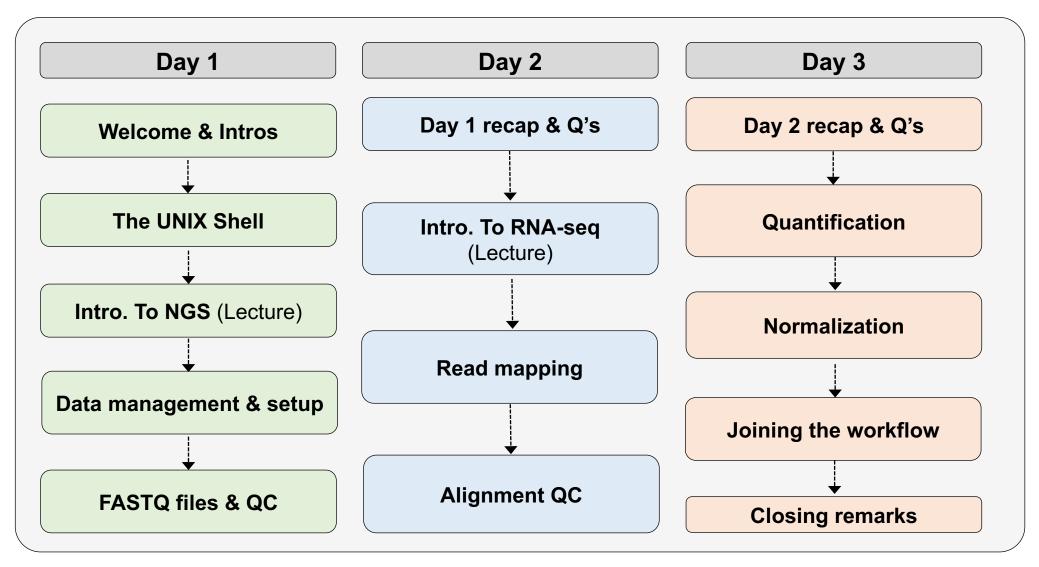
Raw & normalized counts



Differential expression workshop (July)

Workshop outline







Goals of the workshop

- > Develop a working understanding of the analytical workflow for a modern bulk RNA-seq experiment
- > Build a working knowledge of sample preparation considerations for RNA-seq experiments
- > Learn how to process raw NGS data in FASTQ format to generate a gene expression matrix
- ➤ Learn how to perform a detailed quality control analysis

Schedule

Can be found at: https://github.com/Dartmouth-Data-Analytics-Core/RNA-seq-Primary-Data-Analysis-workshop-June-2021/blob/main/schedule.md

> 12pm-5pm each day

Schedule is best guess, and we may deviate from it based on time

➤ If you will be absent for a session, just let us know

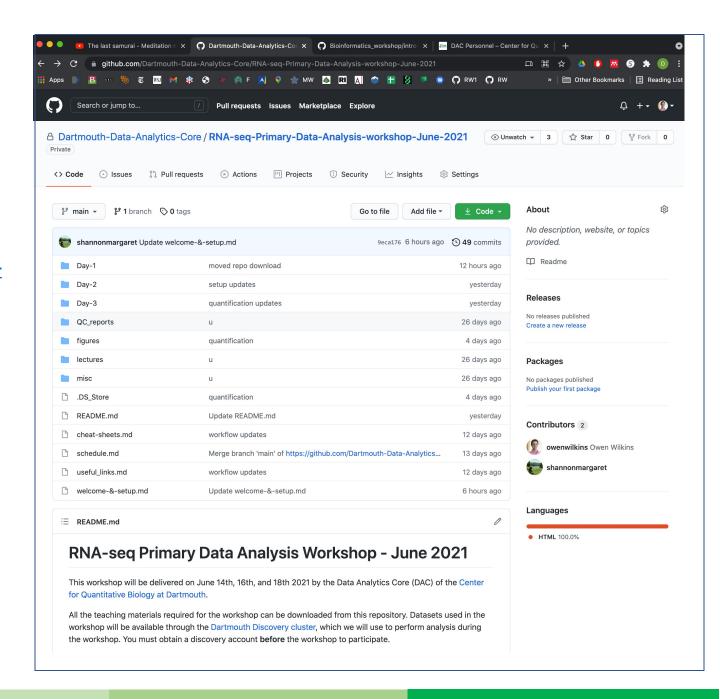
Logistics I

Course materials are all online (and will stay there):

https://github.com/Dartmouth-Data-Analytics-Core/RNA-seq-Primary-Data-Analysis-workshop-June-2021

Lessons, lectures, cheat sheets, etc. are all here

We will be running Bash code from markdowns (.md) into your terminal application

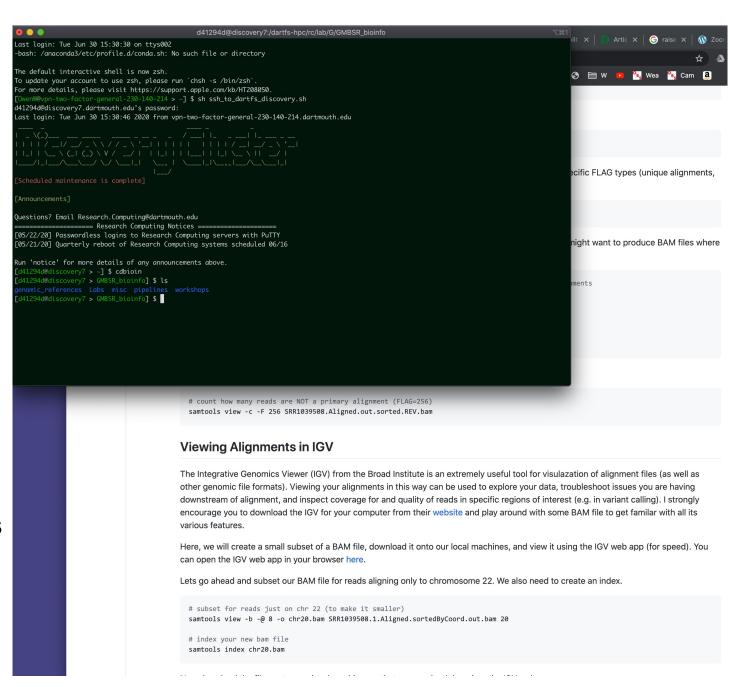


Logistics II

Multiple tabs open

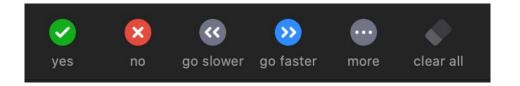
- Terminal window (ssh to discovery7)
- Web browser to GitHub repo

- > If you finish: edit the code, try different options, generate scripts
- Use the cheat sheets!



Logistics III

➤ Use buttons in *Participants* tab in zoom



> You'll be muted, but if you want to ask a question, just raise your hand





- We will use these when we split up to run code independently
- We've tried to pair everyone based on experience
- If your stuck, message us, and we will come help you in your (BR)
- When we are going to move on, breakout rooms will close



Please be courteous on zoom...

How to get help?



> Raise your hand in zoom (bottom right, participants tab)

- > Use the slack channel to message one of us
 - Use the *general* channel if it might benefit everyone
 - Message us directly if its specific



- > If all else fails, email us:
 - DAC: <u>DataAnalyticsCore@groups.dartmouth.edu</u>
 - Shannon Soucy (Shannon.Margaret.Soucy@Dartmouth.edu)
 - Owen Wilkins (<u>omw@Dartmouth.edu</u>)
 - Tim Sullivan (Timothy.J.Sullivan@dartmouth.edu))

Questions after the workshop?

Bioinformatics office hours

Friday 1-2pm (every other week, check calendar: https://sites.dartmouth.edu/cqb/upcoming-events/calendar/

Zoom link: https://dartmouth.zoom.us/s/96998379866

Passcode: bioinfo

> Also, please give us feedback about this workshop, there will be a survey!

And please ask lots of questions!

Questions?

...then.. Introductions!

Name, department/program, research interests, why are you taking the workshop?