Analysis of High Throughput Sequencing RNA-Seq Data

(November 7th -8th); November 14th - November 18th; (November 25th)

Pre-course material (mandatory before Module 2 if Module 1 is skipped)

This will be made available two weeks ahead of the course (October 24th)

Practice

- Get access to a terminal, log in with the user: training (link and password will be provided in updated schedule on Oct 24th)
- An introduction to the unix/linux command line. Do the 10 steps listed on: LinuxCommand.org.
- An introduction to the R language and environment. The website requires to register, but the first lesson of the content linked is accessible for "free": (link will be provided in updated schedule on Oct 24th)
- For a more in depth introduction to R, check the official one:
 "An Introduction to R"

Pre-course material (mandatory)

Read

- o For Day 1
- o DNA Sequencing in the last 40 years
- o The original RNA-Seg publication
- o Best practices for RNA-seq data analysis
- o Guideline for RNA-Seq data analysis
- o For Day 3
- Statistical considerations
- o Normalization Method Comparison
- o For Day 4
- o <u>Differential Expression Methods Comparison</u>
- Study size and its effect on analysis

Watch

o For Day 1:

A video on High Throughput Sequencing - Eric Chow from University California San Franscisco - Duration: **32 minutes**

o For Day 3:

Consideration about Study Design and statistical background - includes subtitles to get you used to my french accent - Duration: **29 minutes**

Module 1 (optional) - November 7th - 8th

Day 1 – 9:00 – 17:00 – Introduction to the Unix Command Line Interface (CLI) 09:00 - 9:30 - Welcome to the course! Daily Learning Objectives 09:30 - 10:30 - CLI Lecture 10:30 - 10:45 - Coffee break 10:45 - 12:00 - Hands-on practical 12:00 - 13:00 - Lunch break 13:00 - 15:00 - Hands-on continued 15:00 - 15:15 - Coffee break 15:15 - 15:30 - "homework" instructions 15:30 - 15:40 - Feedback 15:40 - 17:00 - Self-study and assessment Day 2 – 9:00 – 17:00 – Introduction to the R programming language and the R-Studio environment 09:00 - 9:30 - Day 1 Assessment review / revision session Daily Learning Objectives 09:30 - 10:30 - RStudio environment and R programming lecture 10:30 - 10:45 - Coffee break 10:45 - 12:00 - Hands-on practical 12:00 - 13:00 - Lunch break 13:00 - 15:00 - Hands-on c'ed 15:00 - 15:15 - Coffee break 15:15 - 15:30 - "homework" instructions 15:30 - 15:40 - Feedback 15:40 - 17:00 - Self-study and assessment Module 2 - November 14th - November 18th - 9:00 - 17:00 every day Day 1 - 9:00 - 13:00 - High Throughput Sequencing, 2nd generation 09:00 - 9:30 - Welcome to the course! Daily Learning Objectives 09:30 - 10:30 - High Throughput Sequencing Lecture 10:30 - 10:45 - Coffee break 10:45 - 11:15 - Data pre-processing and analysis workflow 11:15 - 11:30 - Workshop data description 11:30 - 12:40 - Data pre-processing journal club 12:40 - 12:50 - "homework" instructions 12:50 - 13:00 - Feedback In preparation for Day 2 - Self-study and Daily Assessment

Day 2 - 9:00 - 14:00 - Sequencing data pre-processing 09:00 - 10:00- Revision session AND VOTE for Day 5 content! Daily Learning Objectives

10:00 - 10:30 - Sequence format and data pre-processing

10:30 - 10:45 - Coffee break

10:45 - 12:00 - Hands-on practical data pre-processing continued

12:00 - 13:00 - Lunch break

13:00 - 13:40 - Pseudo-alignment Lecture

13:40 - 13:50 - "homework" instructions

13:50 - 14:00 - Feedback

In preparation for Day 3 - Self-study and Daily Assessment

Day 3 - 9:00 - 14:00 - Statistical concepts and Quality Assessment

09:00 - 10:00 - Revision session

Daily Learning Objectives

10:00 - 10:30 - Statistical concerns - Lecture

10:30 - 10:45 - Coffee break

10:45 - 11:30 - Statistical concerns - Lecture

11:30 - 13:40 - Differential Expression analysis setup - until the Exploratory analysis and visualization (included)

12:00 - 13:00 - Lunch break

13:00 - 13:40 - Differential Expression analysis setup - until the Exploratory analysis and visualization (included) c'ed

13:40 - 13:50 - "homework" instructions

13:50 - 14:00 - Feedback

In preparation for Day 4 - Self-study and Daily Assessment

Day 4 - 12:00 - 17:00 - Differential Expression

12:00 - 13:00 - Revision session

Charlotte's extracted code

Revisited code

Revisited code as a html report

Daily Learning Objectives

13:00 - 14:30 - Lecture Differential Expression

14:30 - 14:45 - Coffee break

14:45 - 16:40 - Differential Expression analysis practical

Differential Expression analysis setup - R code

16:40 - 16:50 - "homework" instructions

16:50 - 17:00 - Feedback

In preparation for Day 5 - Self-study and Daily Assessment

Day 5 - 9:00 - 14:00 - Gene Network Inference*

09:00 - 10:00 - Revision session

- Annotated version of Charlotte's DE tutorial

10:00 - 10:30 - Hands-on to kick off network inferences

- Data preparation package

10:30 - 10:45 - Coffee break

10:45 - 11:15 - Gene Network Inference Lecture

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11:15 - 12:00 - Gene Network Inference Practical
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12:00 - 13:00 - Lunch break

13:00 - 13:30 - Gene Network Inference Practical

13:30 - 14:00 - Final Feedback

* This is an exemplary topic; other possible topics are:

Day 5 extra - GSEA Lecture

Day 5 extra - Ensemble GSEA tutorial

Day 5 extra - Machine Learning Lecture

Day 5 extra - Cancer vs. Healthy Machine Learning Prediction

Day 5 extra - Cancer vs. Healthy Machine Learning Classification

Day 5 extra - Single Cell Sequencing lecture

Day 5 extra - Single cell RNA-Seq practical

Day 5 extra - Droplet single cell RNA-Seq practical

Module 3 (optional) - November 25th

Day 1 - 9:00 - 17:00

09:00 - 9:30 - Introduction to the course

9:00 - 10:30 - Interactive sessions to address analysis issues

10:30 - 10:45 - Coffee break

10:45 - 12:00 - Interactive sessions to address analysis issues c'ed

12:00 - 13:00 - Lunch break

13:00 - 14:00 - Presentation preparation

14:00 - 16:00 - Mini-symposium with talks and peer-review

16:00 - 16:45 - Feedback from the trainers on the projects

16:45 - 17:00 - Final Feedback