



OmicsEdu

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Syed Hussain Ather

Olaitan Awe

TJ Butler

Tamiru Denka

Stephen Semick

Wanhua Tang

Background

- Need to educate others
- RNA-Seq is hard
- Create a tutorial others can use
- RNA-Seq and ChIP-Seq analyses

Download raw data
(separate for RNA-Seq and ChIP-Seq)

RNA-Seq

- Preprocessing
 - cutadapt
- Alignment
 - STAR
- Quantification
 - Salmon
 - FeatureCounts
 - HTSeq
- Differential expression
 - DESeq2
- Visualization
 - PCA
 - Boxplots/volcano/heatmaps of DE genes

MultiQC

ChIP-Seq

- Preprocessing
 - picard
 - markduplicates
- Alignment
 - Bowtie
- Peak-calling
 - MACS
- Output comparison
 - Bedtools intersect
- Visualization
 - Heatmaps

Gameplan

- Download everything
- Create workflow overview
 - Run each step individually
 - Put together in jupyter notebook
- Test, test, test
- Create wiki, website