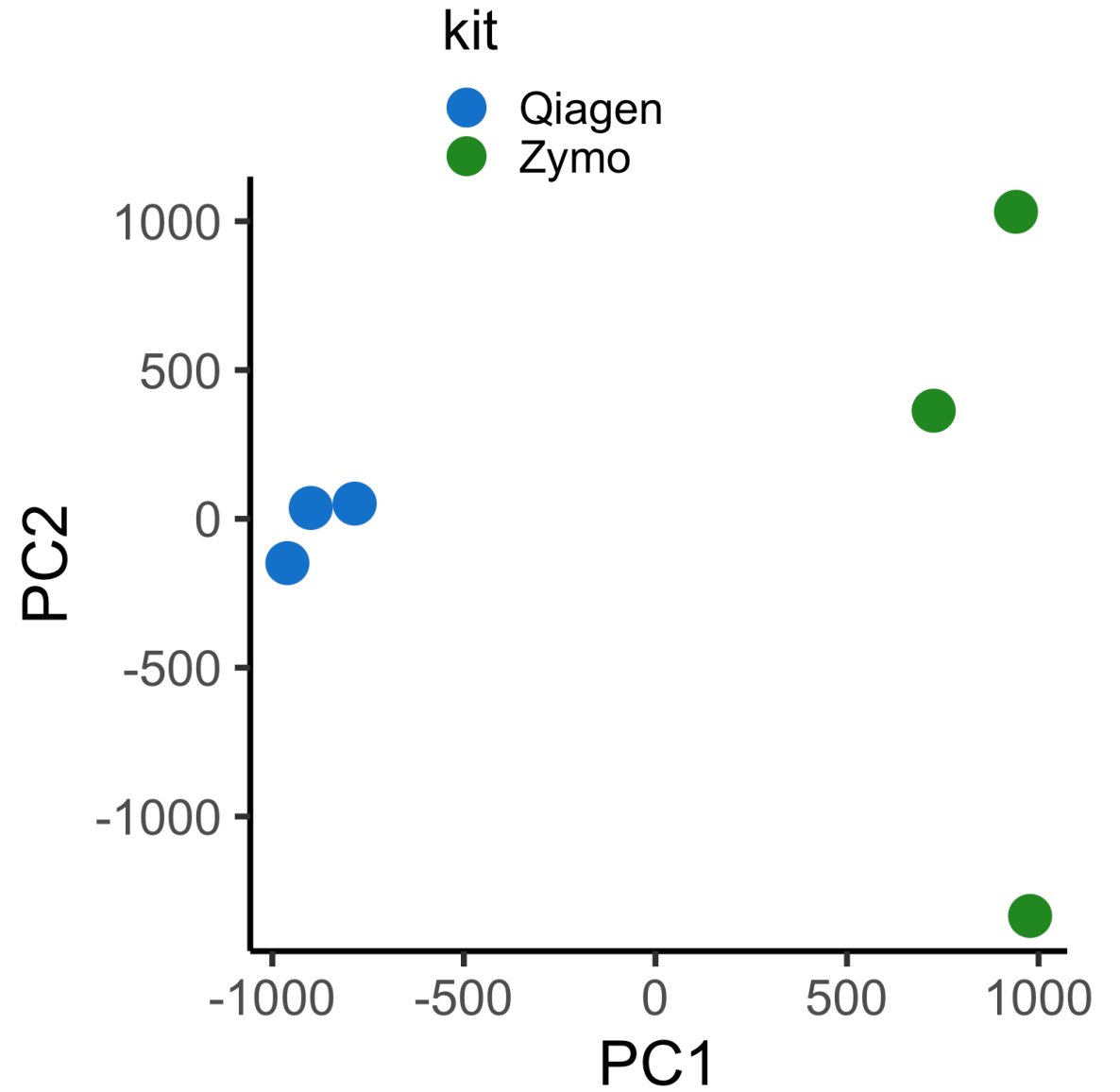


Analyze RNA-seq

2020-07-28

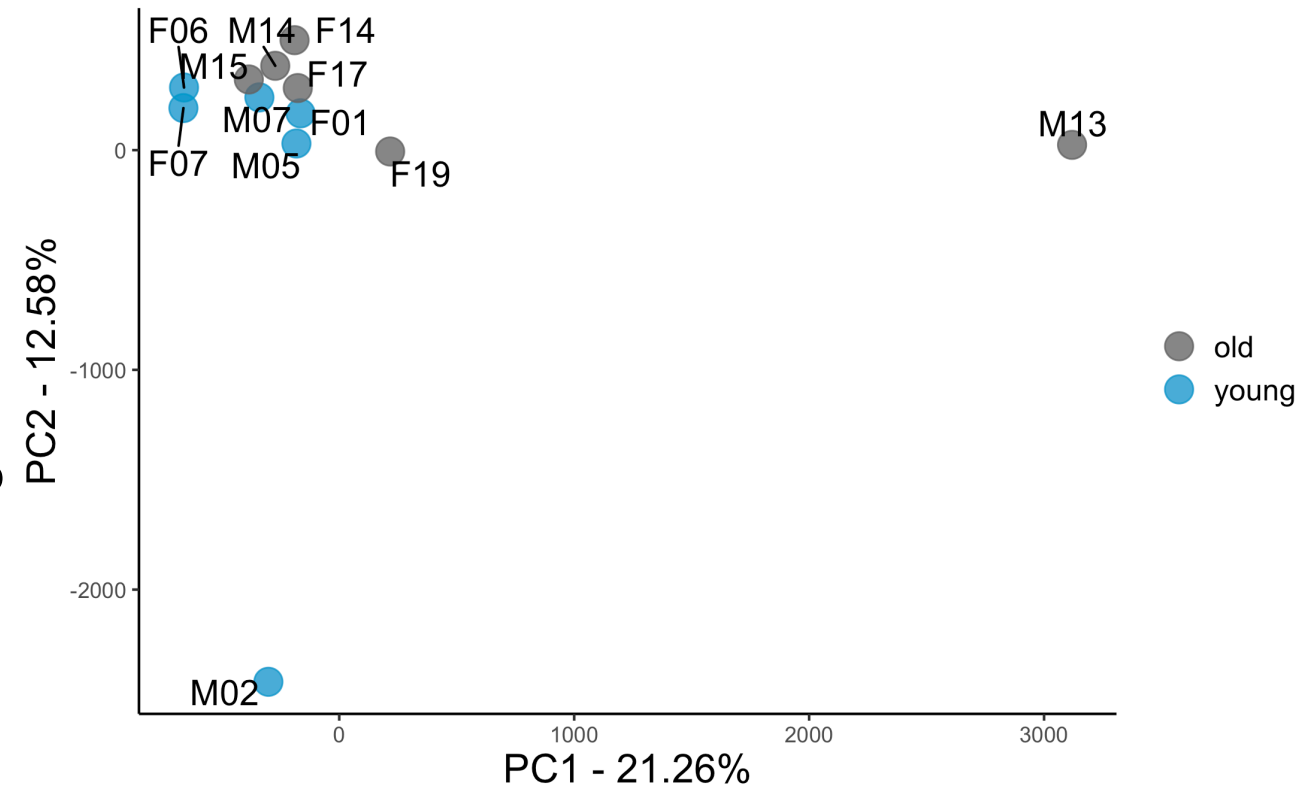
Clustering

- Quality control
 - Is there a batch effect?
 - Is one sample different from the others?
- Do I see expected differences?
- Do I see unexpected differences?



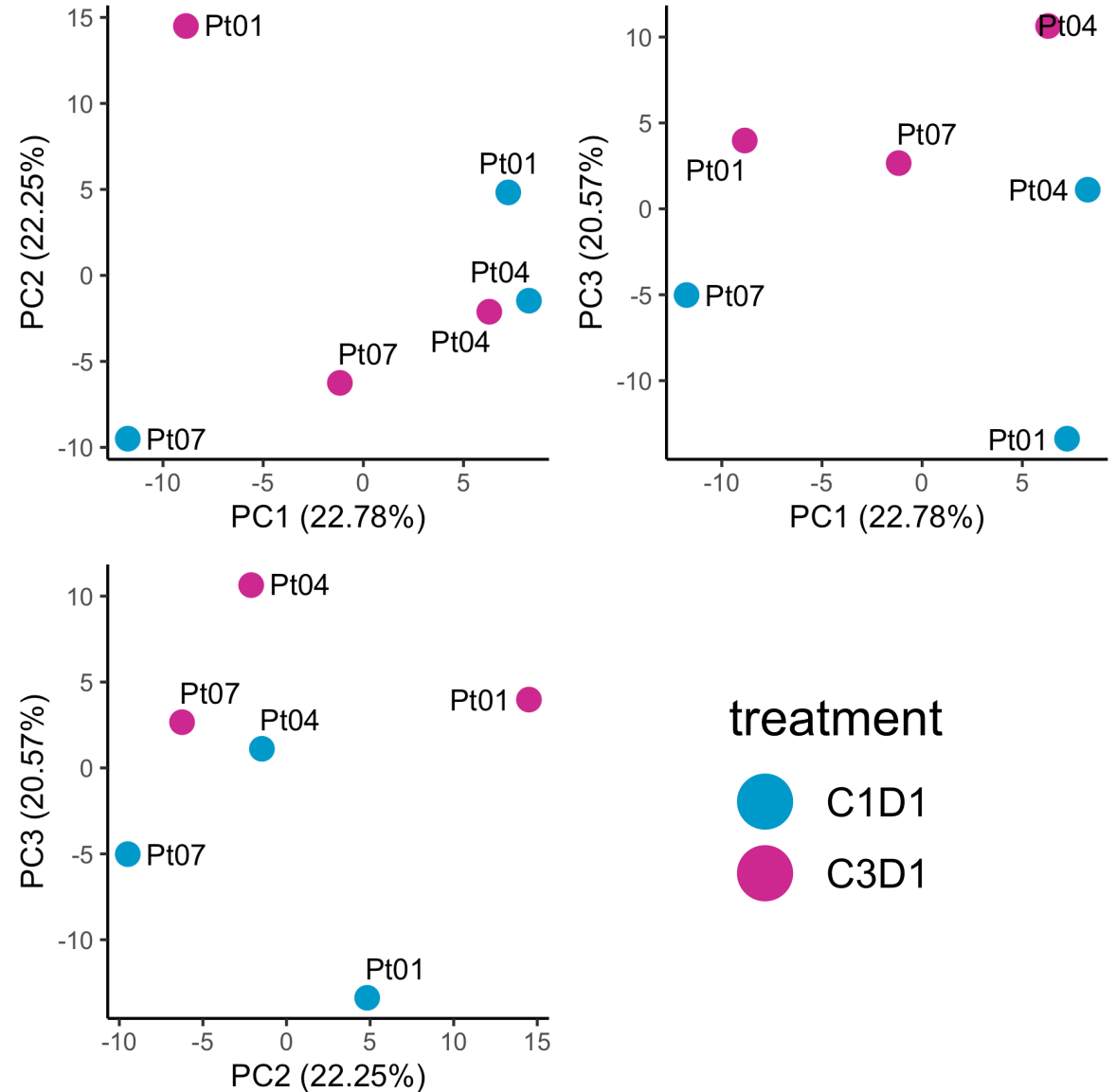
Clustering

- Quality control
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Clustering

- Quality control
 - Is there a batch effect?
 - Is one sample different from the others?
- Do I see expected differences?
- Do I see unexpected differences?



Differential Expression

- Which genes changed expression between control and (whatever variable you manipulated)?
- Test every gene for a difference with a specific type of linear model for RNA-seq
 - log2 fold change between conditions
 - p-value
- Have to correct the results of the tests for multiple testing

Gene	Mean Expression Control	Mean Expression Treatment
Gene A	2	4
Gene B	2	1

Gene A

$$\text{fold change (FC)} = \frac{\text{treatment}}{\text{control}} = \frac{4}{2} = 2$$

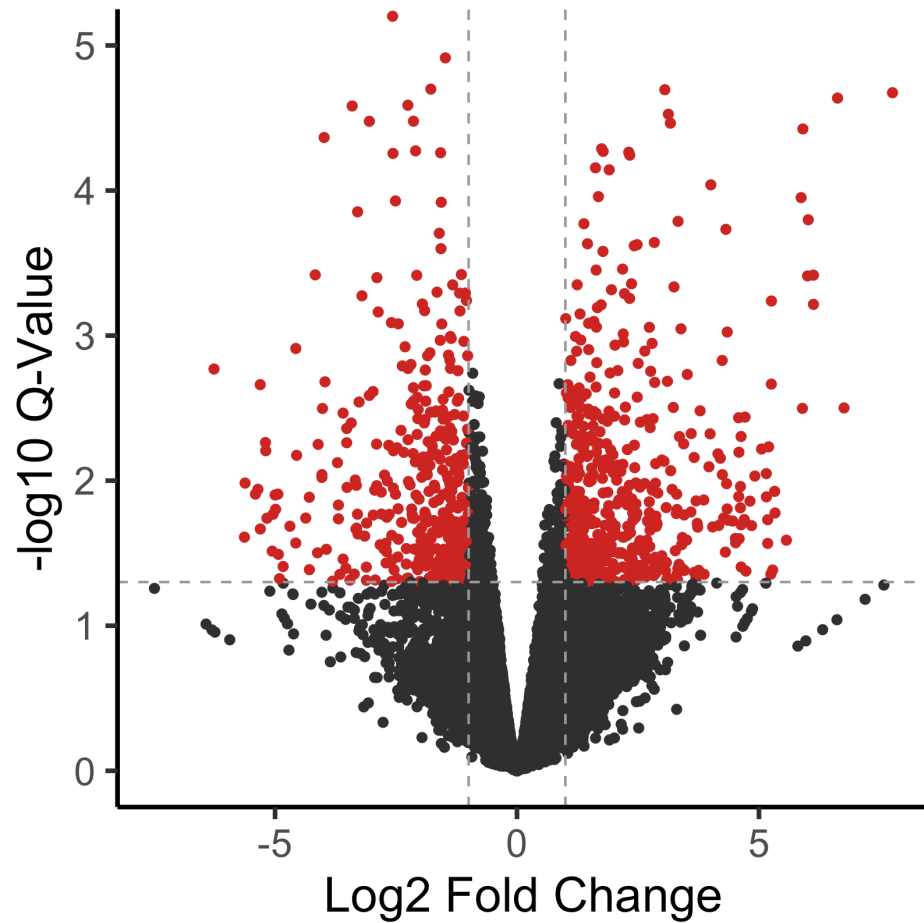
$$\log_2 FC = \log_2 2 = 1$$

Gene B

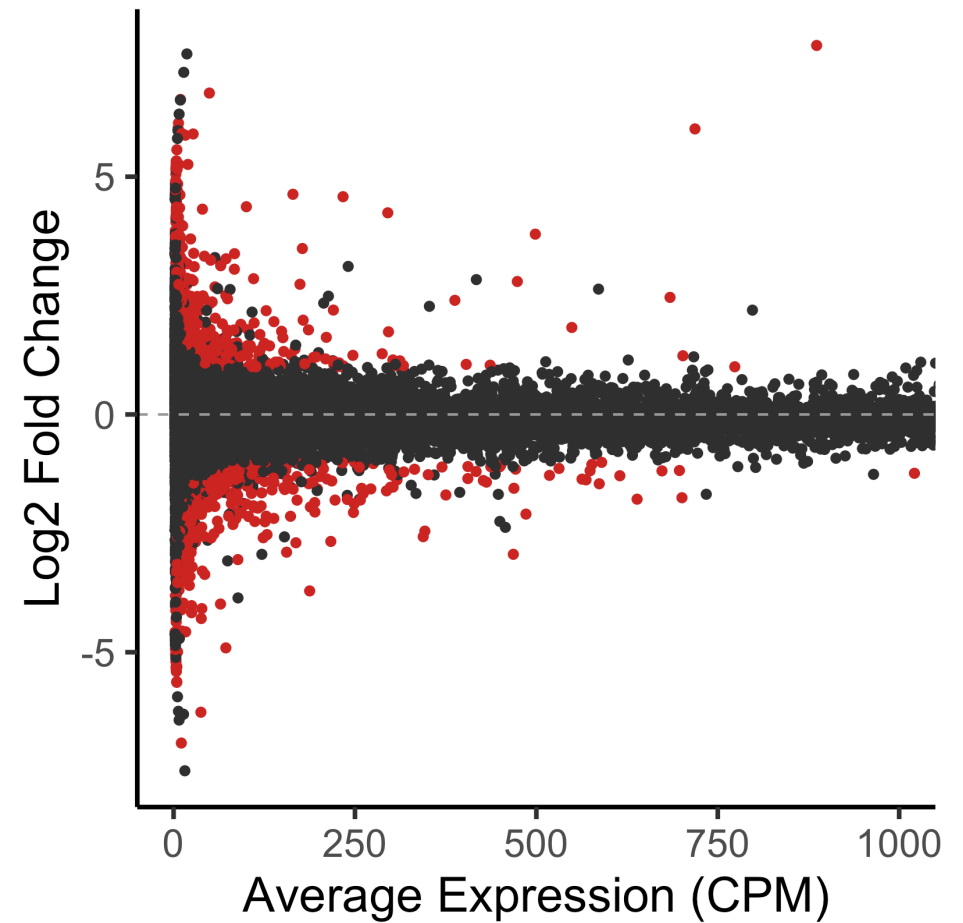
$$\log_2 FC = \log_2 \frac{1}{2} = -1$$

Visualize Results

Volcano Plot



MA Plot



Pathway/Gene Set Enrichment Analysis

- Summarize data
- What sets of genes/signalling pathways/cell circuits have changed?
- Testing for whether you see more genes that belong to the pathway or gene set than expected.

