

Differential gene expression analysis : Example of RNAseq

- Use statistics to compare 2 groups:

For each gene i , is there a **significant difference** in mean expression between control and patients?

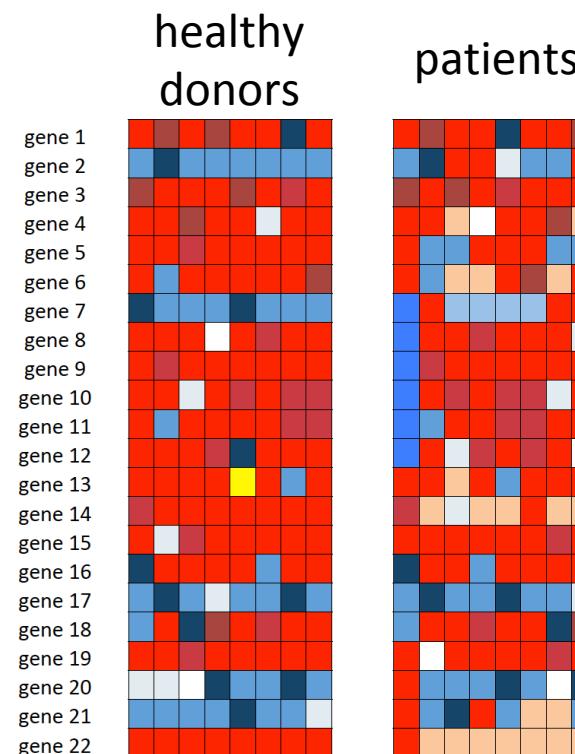
- T-test:

H_0 : Healthy donors and patients have similar gene i expression

$$H_{0i} : \pi_{i1} = \pi_{i2}$$

H_1 : Healthy donors and patients don't have a similar gene i expression

$$H_{1i} : \pi_{i1} \neq \pi_{i2}$$



T-test in R

```
> t.test(grp1, grp2, paired = F)
```

Welch Two Sample t-test

data: grp1 and grp2

t = -6.3689, df = 8.9195, p-value = 0.0001352

alternative hypothesis: true difference in means is not equal to 0

95 percent confidence interval:

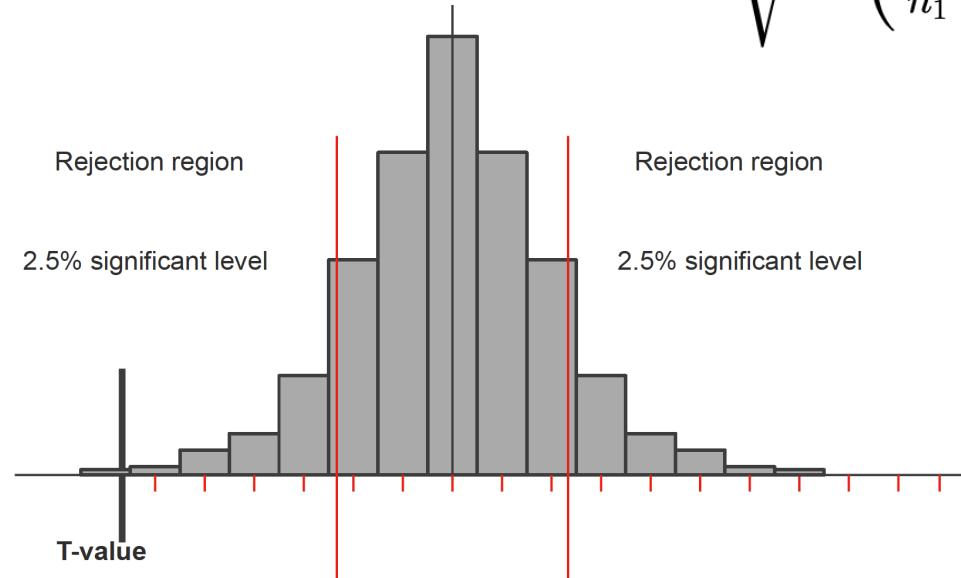
-8.908753 -4.234104

sample estimates:

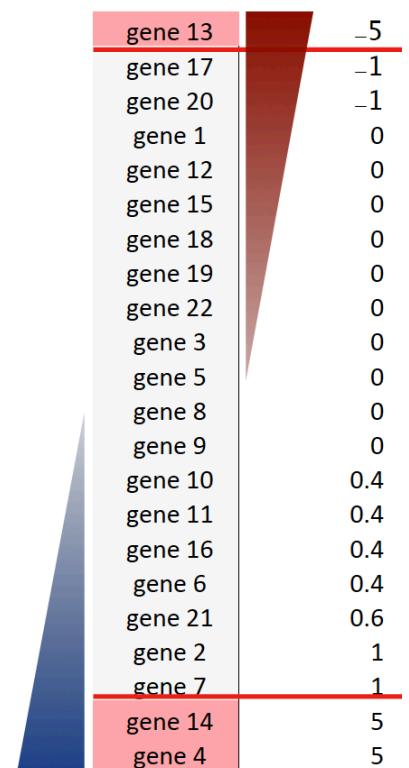
mean of x mean of y

6.00000 12.57143

$$t = \frac{\bar{x}_1 - \bar{x}_2}{\sqrt{s^2 \left(\frac{1}{n_1} + \frac{1}{n_2} \right)}}$$



sort based
on T-statistic



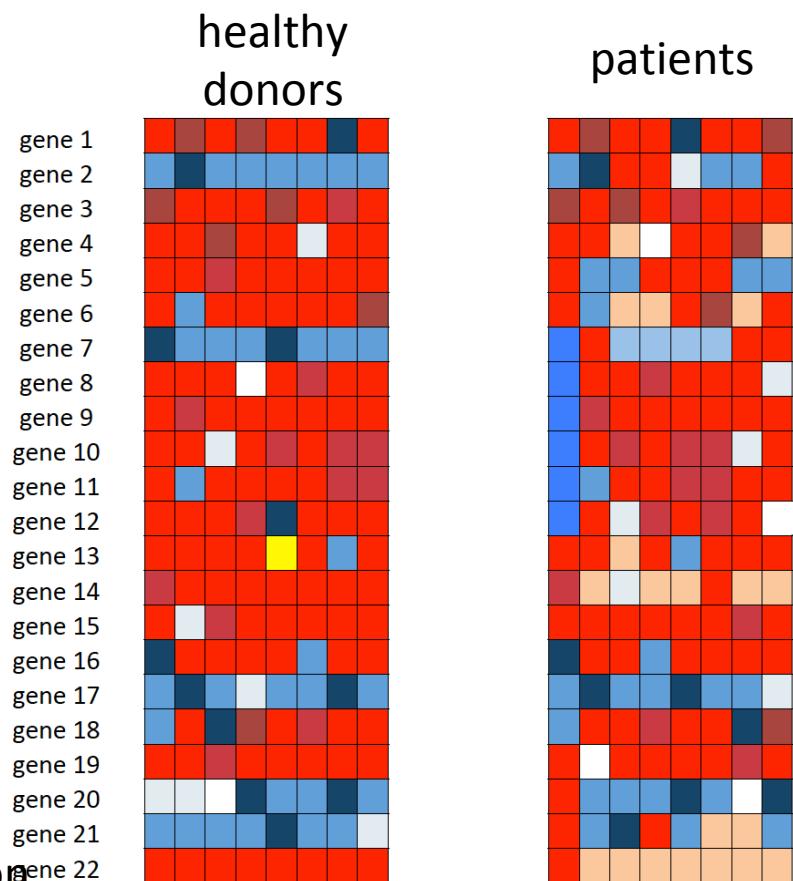
Differential gene expression analysis using R

- Bioconductor

<https://bioconductor.org/>

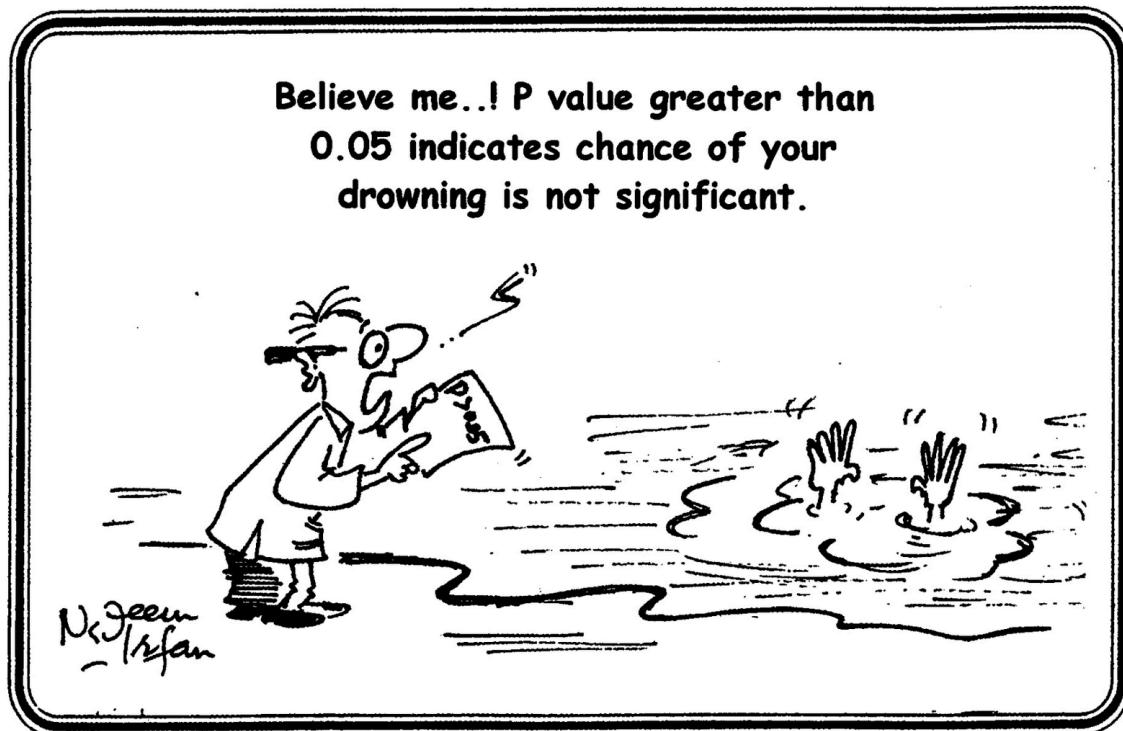
- Several packages :
 - limma: t-test
 - DESeq2: Wald test
 - edgeR: exact test

All allow for simultaneous statistical tests
for every gene, together with p-value calculation



What does $p < 0.05$ mean?

- It implies that it is acceptable to have a 5% probability to incorrectly reject the null hypothesis while it is correct.
- It means that if we repeat an experiment 20 times, we would reject the null hypothesis once because of random error.



P-value adjustment: what is it?

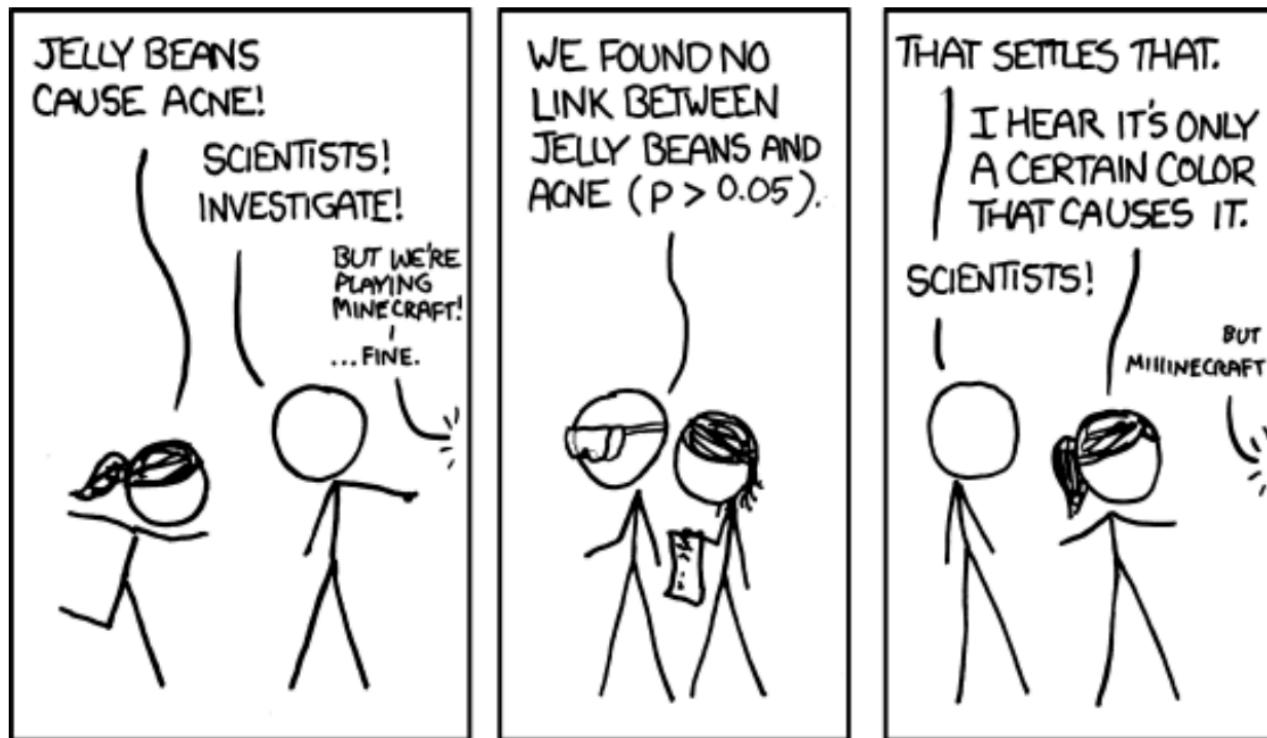


Photo by Patrick Fore on Unsplash

Cartoon: <https://xkcd.com/882/>

Paper on p-value adjustment: <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC6099145/>

WE FOUND NO
LINK BETWEEN
PURPLE JELLY
BEANS AND ACNE
($P > 0.05$).


WE FOUND NO
LINK BETWEEN
BROWN JELLY
BEANS AND ACNE
($P > 0.05$).


WE FOUND NO
LINK BETWEEN
PINK JELLY
BEANS AND ACNE
($P > 0.05$).


WE FOUND NO
LINK BETWEEN
BLUE JELLY
BEANS AND ACNE
($P > 0.05$).


WE FOUND NO
LINK BETWEEN
TEAL JELLY
BEANS AND ACNE
($P > 0.05$).


WE FOUND NO
LINK BETWEEN
SALMON JELLY
BEANS AND ACNE
($P > 0.05$).


WE FOUND NO
LINK BETWEEN
RED JELLY
BEANS AND ACNE
($P > 0.05$).


WE FOUND NO
LINK BETWEEN
TURQUOISE JELLY
BEANS AND ACNE
($P > 0.05$).


WE FOUND NO
LINK BETWEEN
MAGENTA JELLY
BEANS AND ACNE
($P > 0.05$).


WE FOUND NO
LINK BETWEEN
YELLOW JELLY
BEANS AND ACNE
($P > 0.05$).


WE FOUND NO
LINK BETWEEN
GREY JELLY
BEANS AND ACNE
($P > 0.05$).


WE FOUND NO
LINK BETWEEN
TAN JELLY
BEANS AND ACNE
($P > 0.05$).


WE FOUND NO
LINK BETWEEN
CYAN JELLY
BEANS AND ACNE
($P > 0.05$).


WE FOUND A
LINK BETWEEN
GREEN JELLY
BEANS AND ACNE
($P < 0.05$).
WHAOA!


WE FOUND NO
LINK BETWEEN
MAUVE JELLY
BEANS AND ACNE
($P > 0.05$).

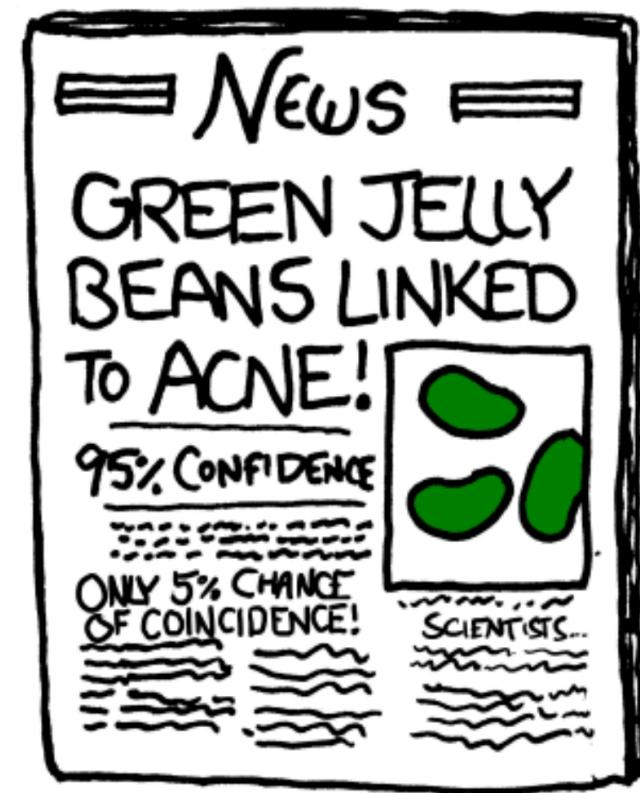

WE FOUND NO
LINK BETWEEN
BEIGE JELLY
BEANS AND ACNE
($P > 0.05$).


WE FOUND NO
LINK BETWEEN
LILAC JELLY
BEANS AND ACNE
($P > 0.05$).


WE FOUND NO
LINK BETWEEN
BLACK JELLY
BEANS AND ACNE
($P > 0.05$).


WE FOUND NO
LINK BETWEEN
PEACH JELLY
BEANS AND ACNE
($P > 0.05$).


WE FOUND NO
LINK BETWEEN
ORANGE JELLY
BEANS AND ACNE
($P > 0.05$).

Methods of p-value adjustment

- **Bonferroni:** the alpha level is divided by the total number of tests
 - if we run $k=20$ tests:
 $0.05/k = 0.05/20=0.0025$
- Good for small number of tests but too conservative for thousands of genes
- **Benjamini-Hochberg procedure (BH, decreases the FDR)**
 - Rank the p-values from smallest to largest, adjust less and less as the p-values get larger:
 $p\text{-value}_1 * (n/1)$
 $p\text{-value}_2 * (n/2)$
...
 $p\text{-value}_k * (n/k) = p\text{-value}_k * 1$
- n = total number of p-values (genes)
 k = rank number of each p-value

<https://www.youtube.com/watch?v=rZKa4tW2Nks>