

The Count Matrix

| | Condition 1 | | | Condition 2 | | |
|-------|-------------|------|------|-------------|------|------|
| | Rep1 | Rep2 | Rep3 | Rep1 | Rep2 | Rep3 |
| Gene1 | 72.7 | 75.7 | 66.3 | 55.0 | 51.2 | 35.9 |
| Gene2 | 81.3 | 77.4 | 71.1 | 58.1 | 47.9 | 32.8 |
| Gene3 | 80.2 | 75.3 | 70.0 | 57.1 | 51.9 | 43.3 |
| Gene4 | 78.8 | 76.7 | 68.8 | 58.0 | 43.9 | 41.5 |
| Gene5 | 79.8 | 74.6 | 67.9 | 60.2 | 45.3 | 38.5 |
| | 76.1 | 74.5 | 69.7 | 59.6 | 45.3 | 40.5 |
| | 78.8 | 79.0 | 74.5 | 58.0 | 52.8 | 50.8 |
| | 78.7 | 79.2 | 71.8 | 58.8 | 49.8 | 38.3 |
| | 76.8 | 74.0 | 70.5 | 64.1 | 46.6 | 33.4 |

Hands-on!

- Getting familiar with the tidyverse
- Exploring RNAseq Counts
- Differential gene expression analysis
- Visualisation techniques
- *Functional enrichment analysis*

https://hcliedtke.github.io/UAM23_rnaseq/