

- Command
- ▭ Output (example report.ppt)

Output1

Global view (Three line table)

Six columns:

- 1) Sample name
- 2) Raw reads
- 3) Filtered reads
- 4) Unique mapped reads
- 5) Removed duplicate reads
- 6) Gene reads

Output3

- 1) Gene Expression (TPM)
- 2) PCA plot
- 3) DEG (Differential Expression Gene)
- 4) volcano plot
- 5) DEG heatmap
- 6) DEG GO enrichment
- 7) DEG KEGG enrichment

