

- ▼  test
 - ▶  fastqc
 - ▶  fastqc_after_trimming
 - ▼  genome
 - ▶  alignmentQC
 - ▶  countFile
 - ▼  dea
 - ▶  countGroup
 - ▶  DEA
 - ▶  visualization
 - ▶  report_align_count_data
 -  report_align_count.html
 - ▼  trans
 - ▼  dea
 - ▶  countGroup
 - ▼  DEA
 - ▶  gene-level
 - ▶  transcript-level
 - ▶  visualization
 - ▶  quant
 - ▶  report_quantify_data
 -  report_quantify.html
 - ▶  tpmFile