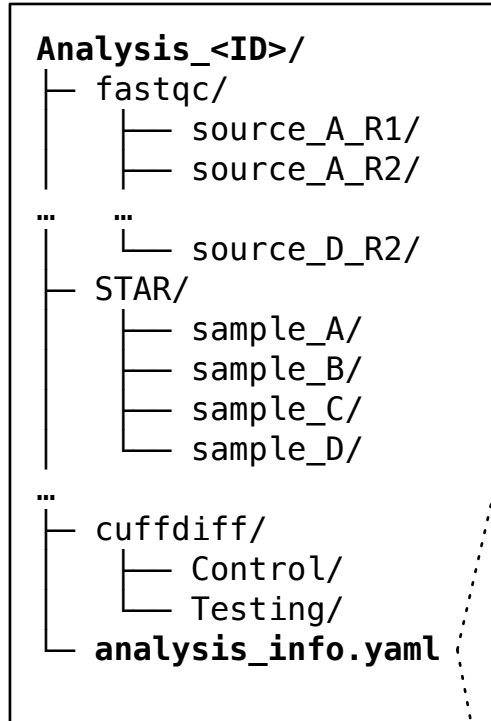


*Analysis result
of Cufflinks pipeline*



*Analysis information
in YAML format*

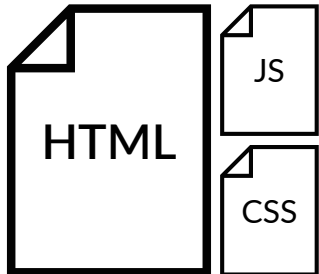
```
conditions:
- Control:
  - A:
    - A_R1.fastq
    - A_R2.fastq
  - B:
    - B_R1.fastq
    - B_R2.fastq
- Testing:
  - C:
    .....
  - D:
    .....

data_sources:
- A_R1.fastq:
  path: 1/A_R1.fastq
  type: FASTQ
  strand: R1
- A_R2.fastq:
  path: 1/A_R2.fastq
  type: FASTQ
  strand: R2
.....

parameters:
STAR:
  runRNGseed: 777

cuffdiff:
  FDR: 0.05
  library-type:
    fr-unstranded
.....
```

*Report templates
for Cufflinks pipeline*



Analysis Results

BCReport

Result Parser

Report Renderer

Analysis Report

Report Templates