

# The three data tables in genomics

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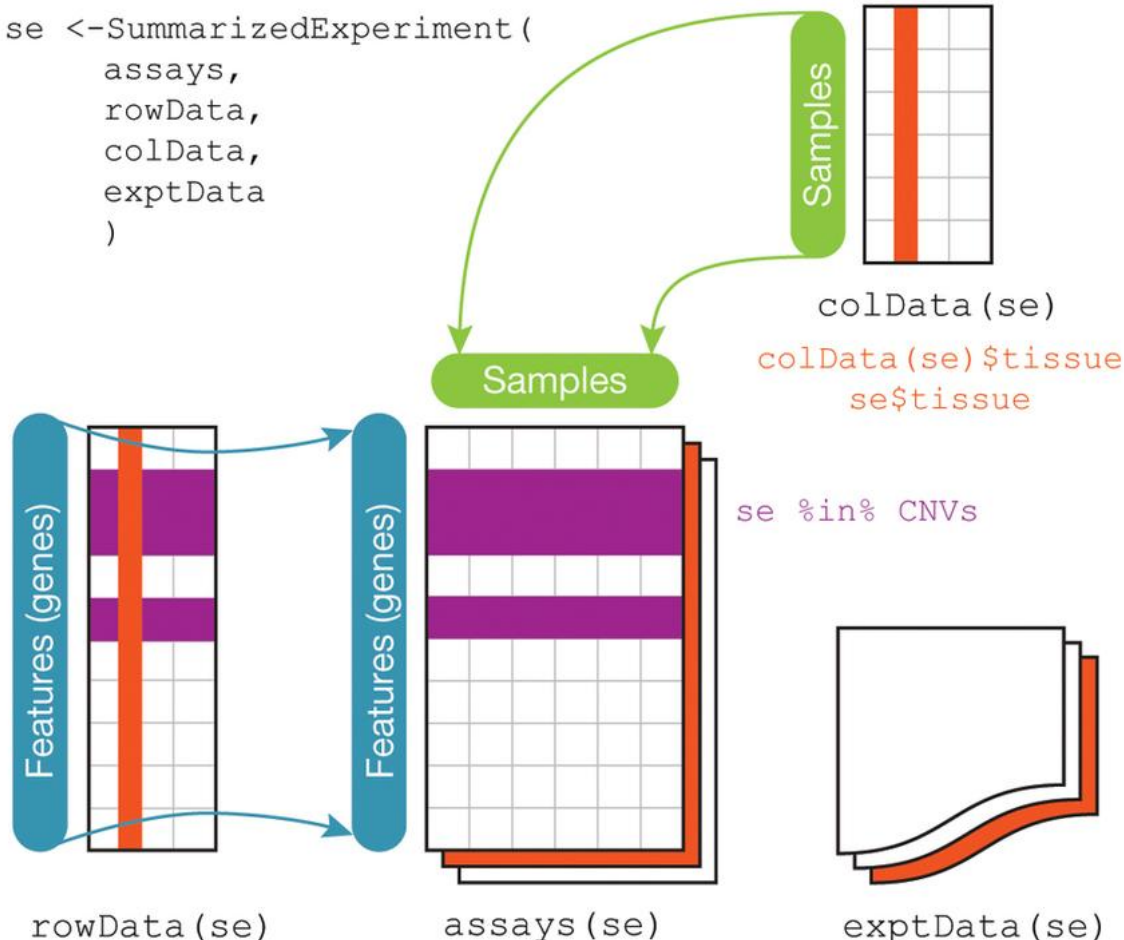
# The three tables

Genomic data

Phenotype data

Feature data

```
se <- SummarizedExperiment (  
  assays,  
  rowData,  
  colData,  
  exptData  
)
```



```
rowData (se)$entrezId assays (se)$count exptData (se)$projectId
```