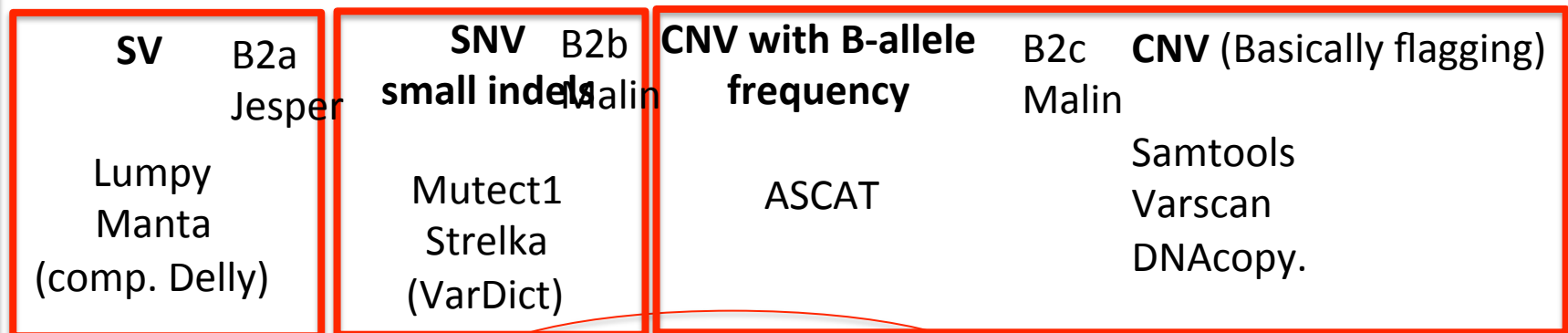
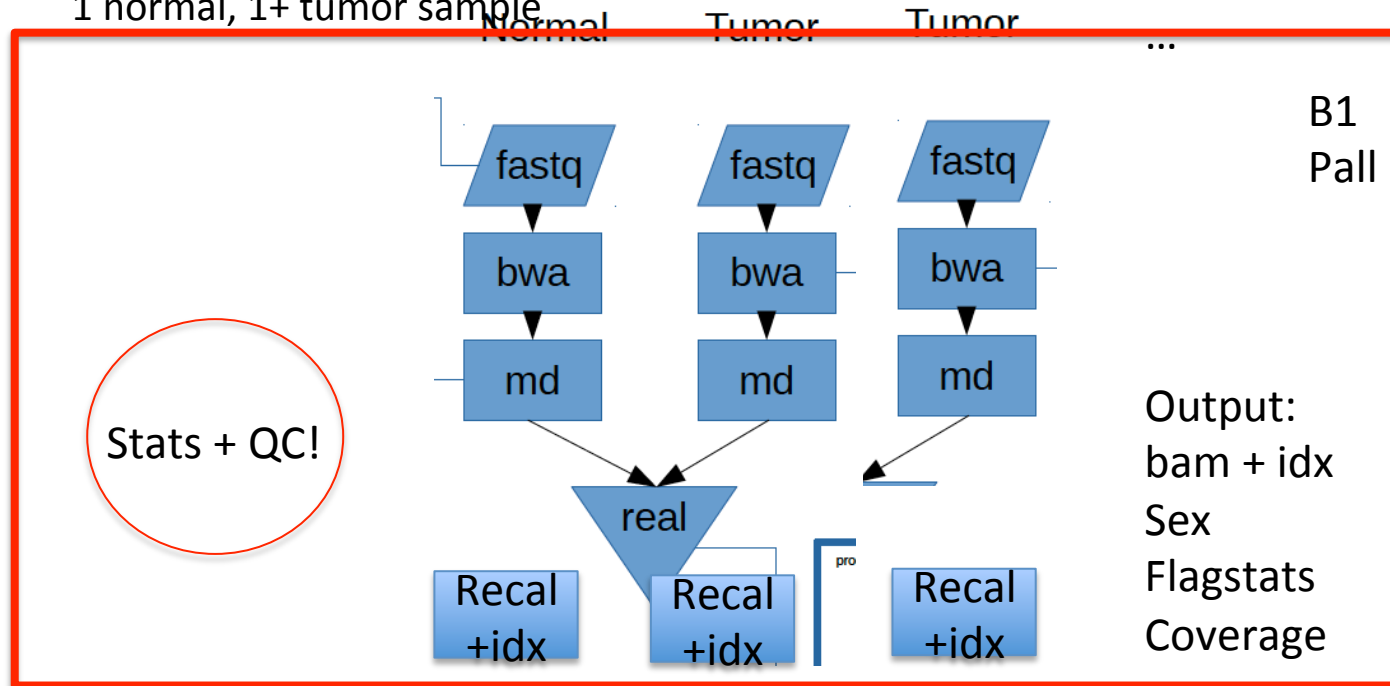


Metadata: (sex)  
>1 fastq files per sample  
1 normal, 1+ tumor sample

Phase1



Annotation and ranking

RNA, methylation, ...

Visualization

Relapse / Tumor evolution

## Annotation and ranking

vep  
SNPeff  
Frequency of variant in pop

Ranking  
Variant qual values? (Read depth, F/R reads..)

B3  
Szilva  
Maxime  
Teresita