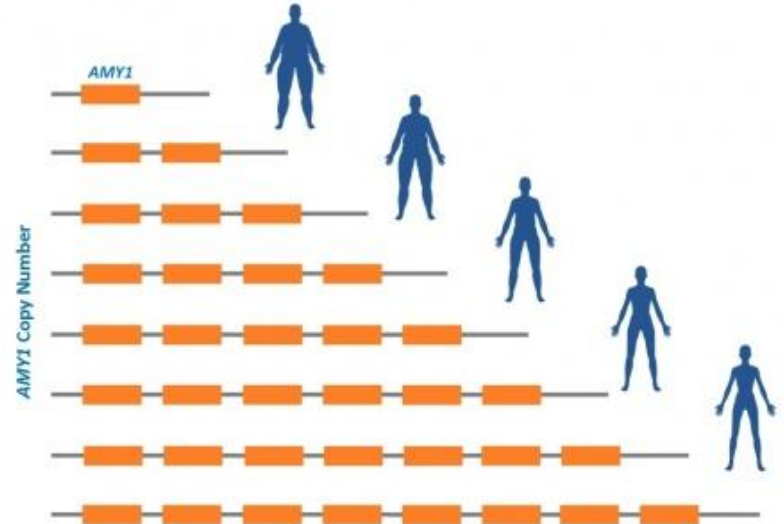
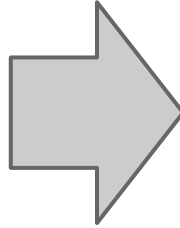


What is computational biology software?

Steven Salzberg

Computational biology software
analysis of genomic data
visualization of results
statistical analysis

[illegible]

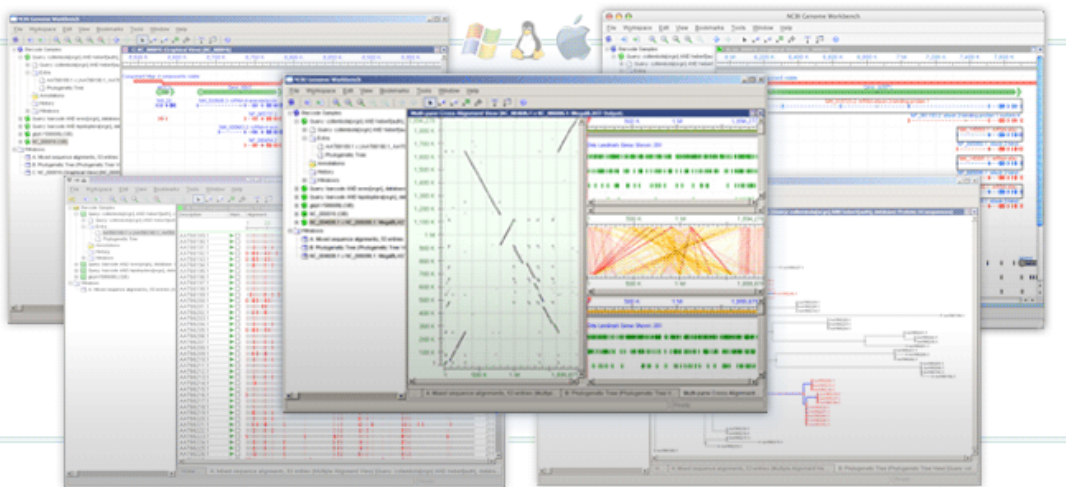
Analysis “pipelines”

DNA

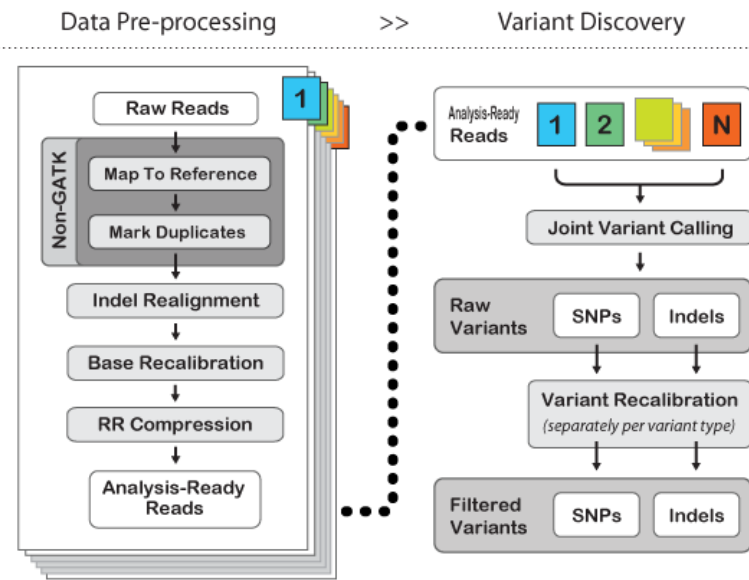
Brilliant
insight



100s of programs and pipelines



NCBI Genome Workbench

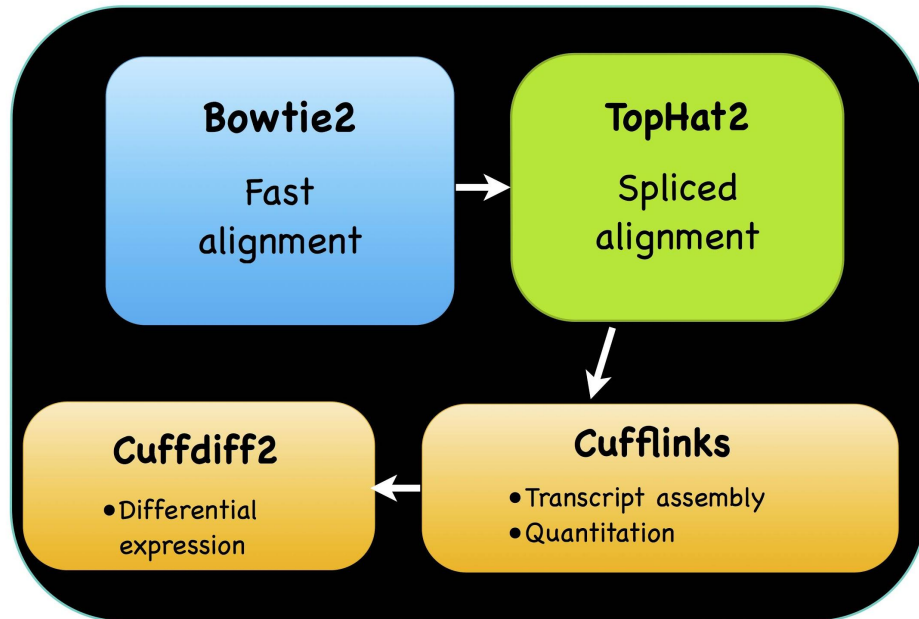


Genetic variant calling

An RNA-seq pipeline



the “Tuxedo” tools



Tuxedo turns reads into genes...

Splicing structure of gene "X"

Relative abundance
of isoforms

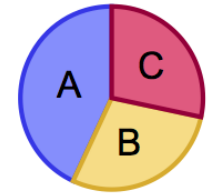
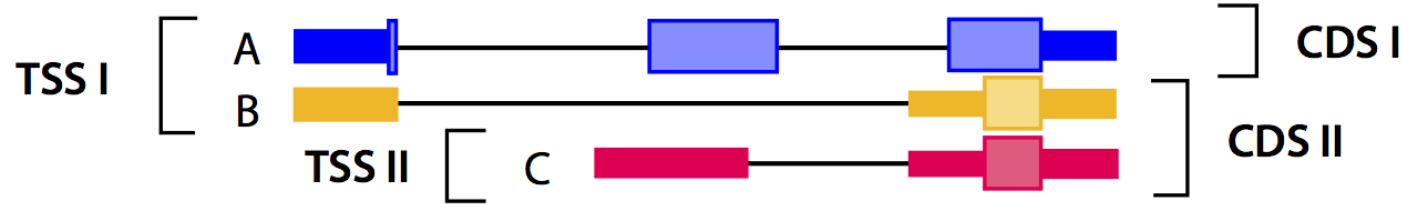
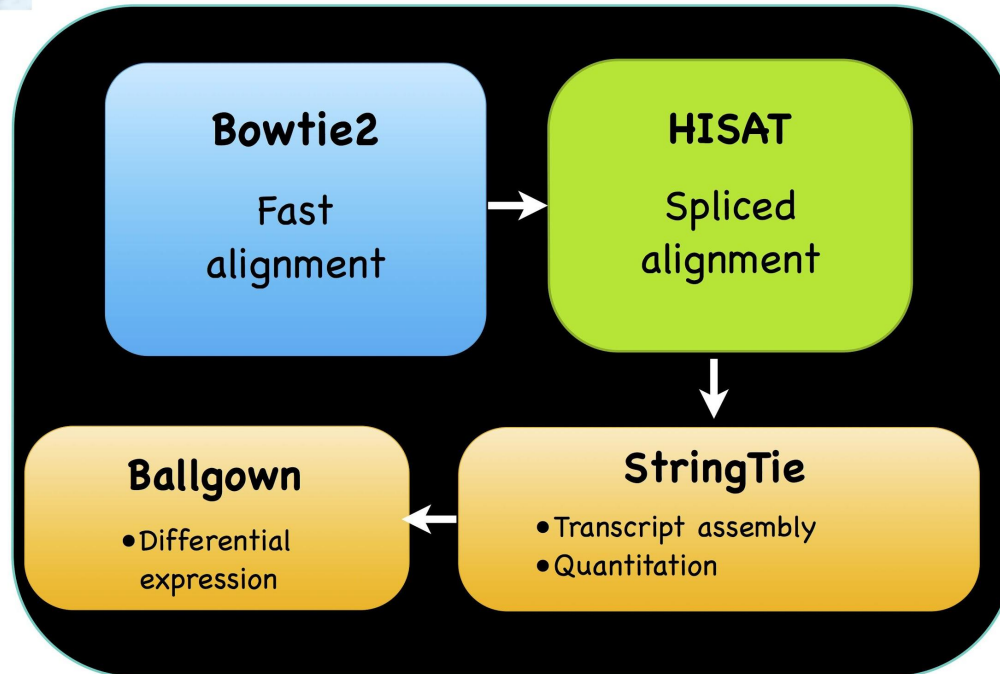


Figure from Trapnell et al (2012),
Nature Protocols 7:3, 562

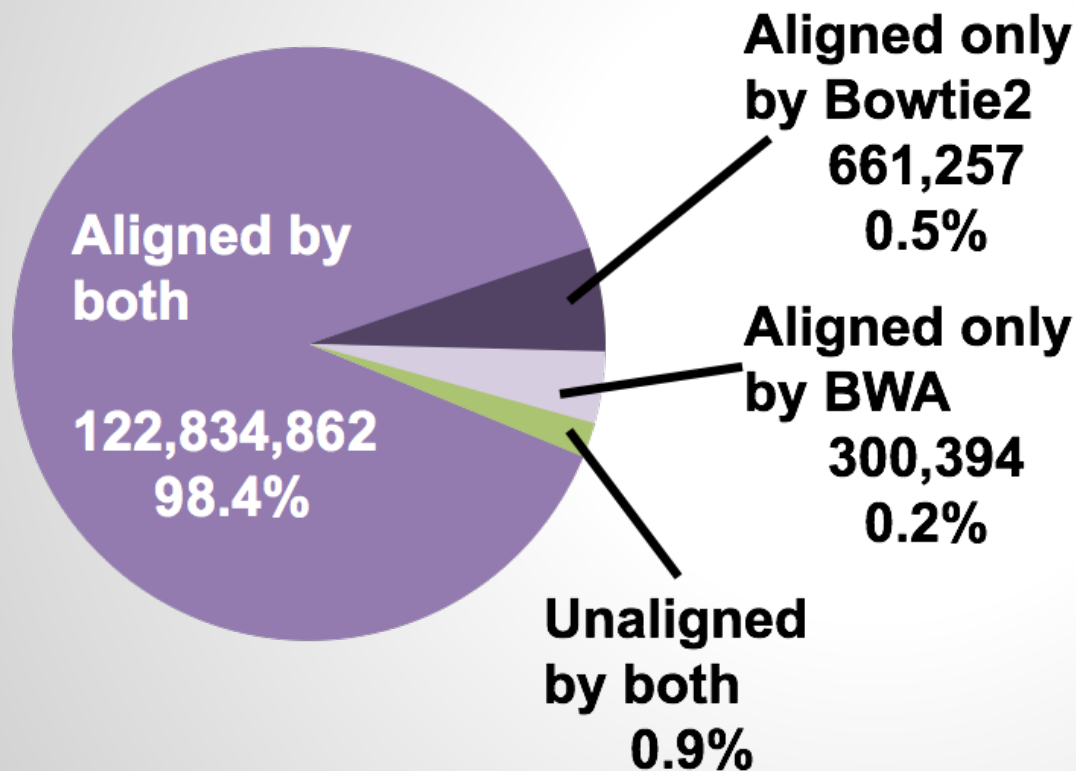
Pipelines change - keep up!



the next-generation
"Tuxedo" tools



Does software choice matter?



Fast-changing software

- Technology is changing
- Software must keep up
- So must we!

