What is computational biology software?

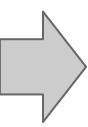
Steven Salzberg

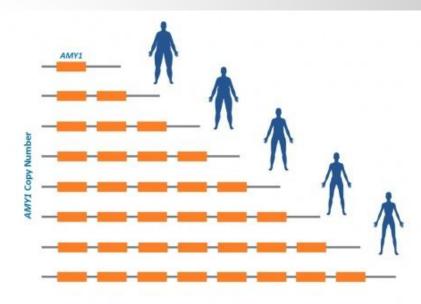


Computational biology software analysis of genomic data visualization of results statistical analysis

Raw data must be transformed

ATCGATCGATGCATCGATCGATCG





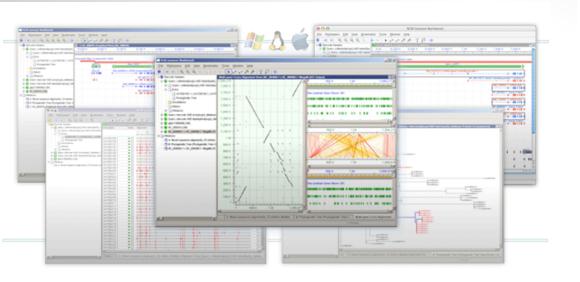
Analysis "pipelines"

Brilliant insight

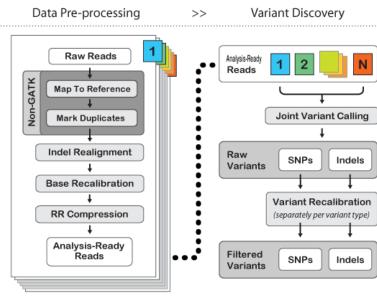
DNA



100s of programs and pipelines



NCBI Genome Workbench

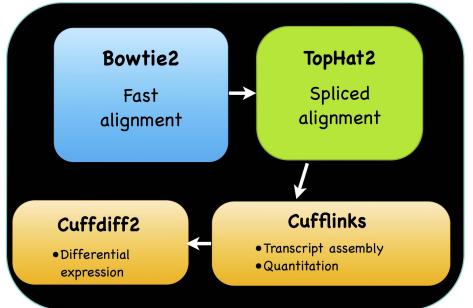


Genetic variant calling

An RNA-seq pipeline



the "Tuxedo" tools



Langmead et al. 2009; Langmead and Salzberg, 2012; Trapnell, Pachter, and Salzberg, 2009; Trapnell et al., 2010; Kim et al., 2013

Tuxedo turns reads into genes...

Splicing structure of gene "X"

Relative abundance of isoforms



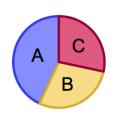
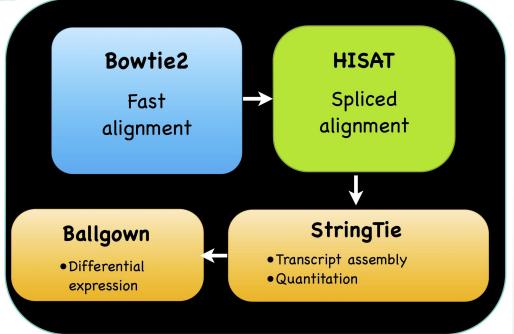


Figure from Trapnell et al (2012), *Nature Protocols* 7:3, 562

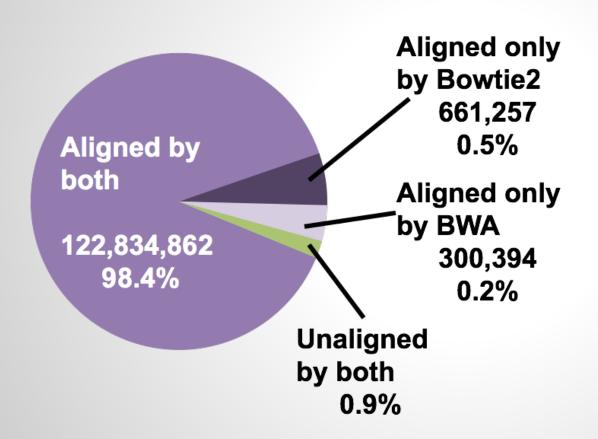
Pipelines change - keep up!



the next-generation "Tuxedo" tools



Does software choice matter?



Fast-changing software

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

