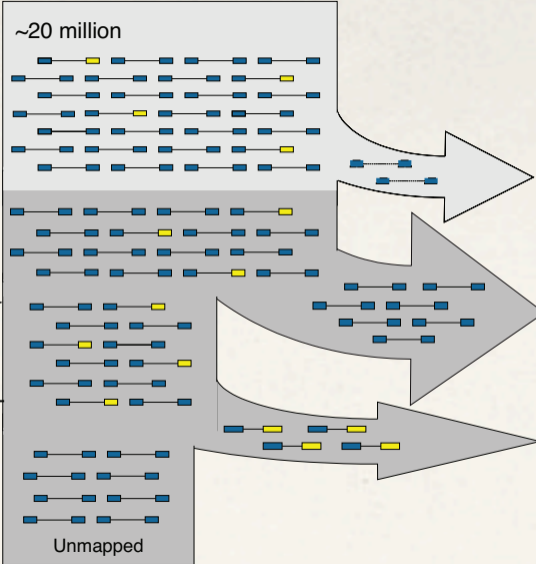


Fusion Detection Module

reference human genome sequence + annotation



Quality control
~0.5 million

Mapping to the same gene
~10 million

Mapping to *different* genes
~ 1000 (1K)

