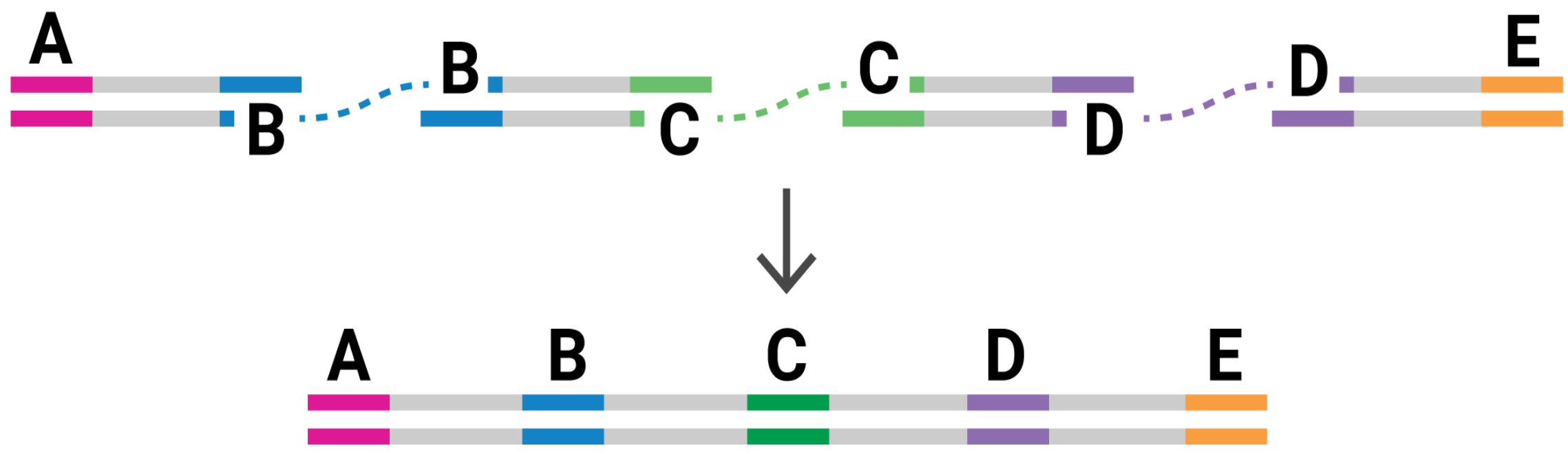


PacBio



**Deconcat
reads**



**Demultiplex
reads**



**Length filter +
FastQC**



**Align conserved
regions**



mcfrieth/last

**Split reads
(UMIs / variants)**

Cluster UMIs



gui11aume/starcode

Cluster variants



mcfrieth/lamassemble

**Combine
consensus data**