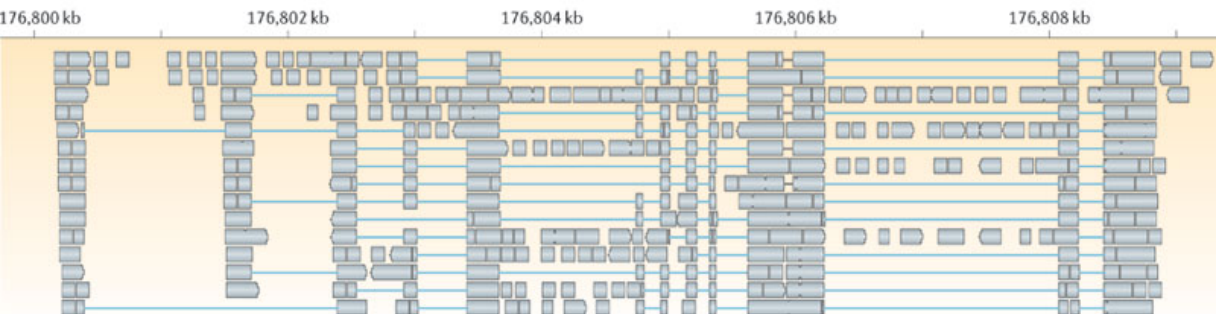
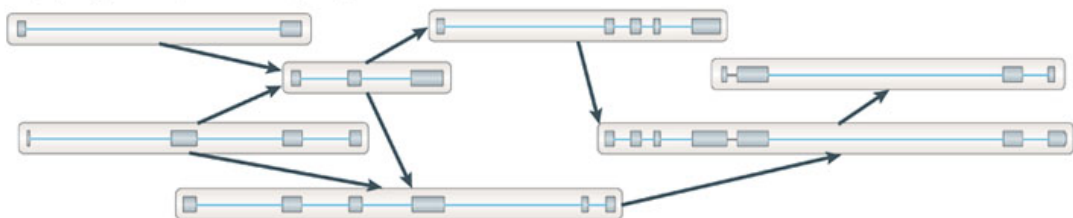


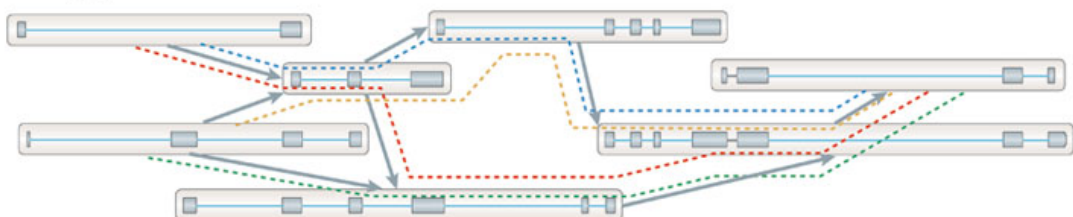
a Splice-align reads to the genome



b Build a graph representing alternative splicing events



c Traverse the graph to assemble variants



d Assembled isoforms

