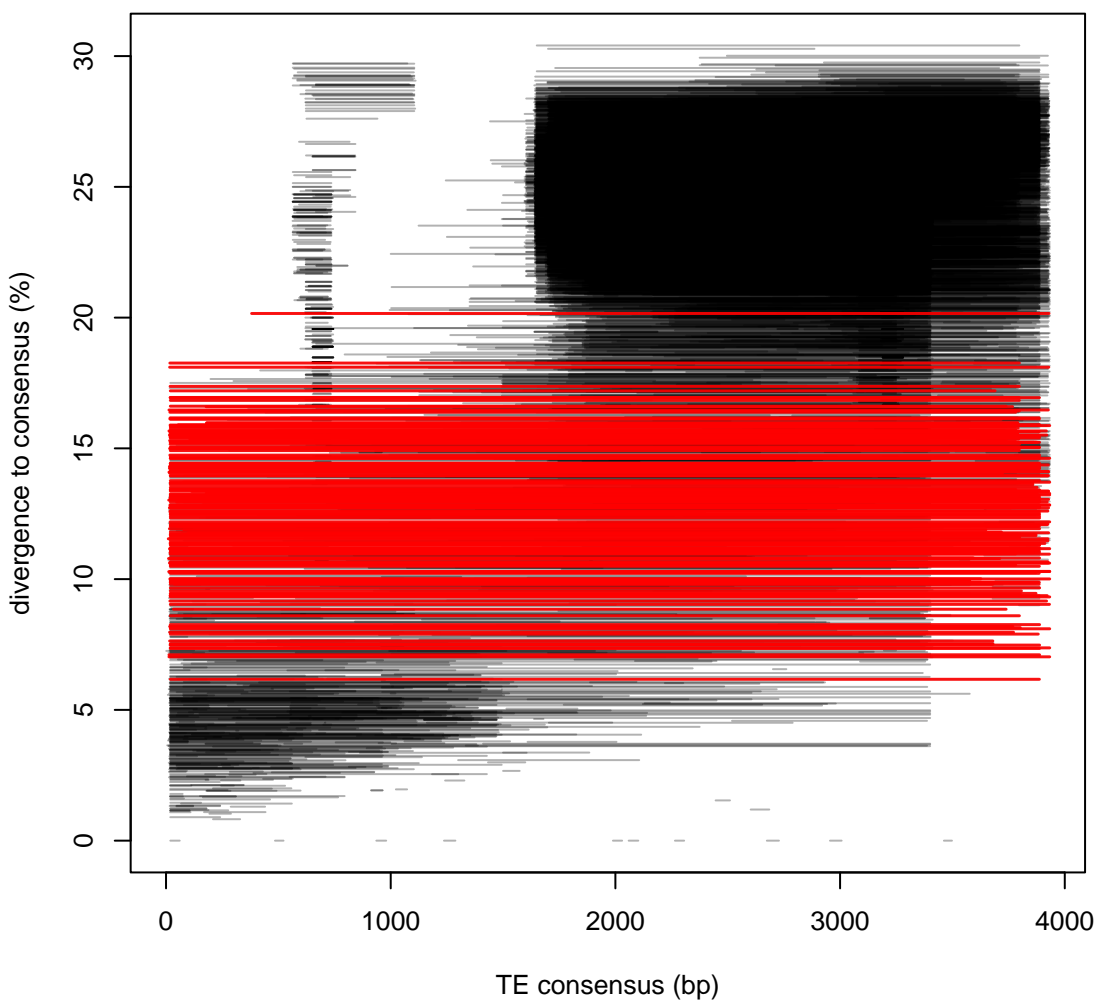
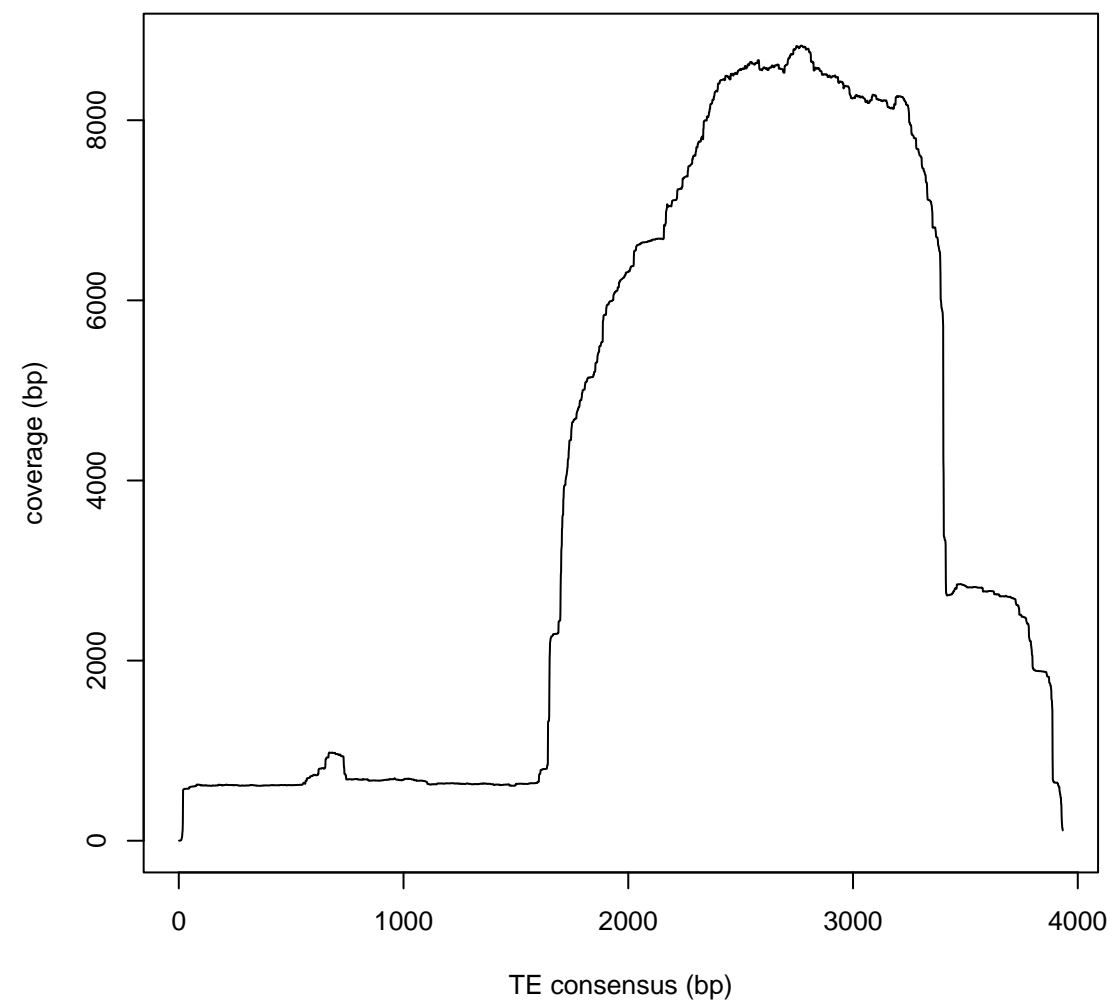


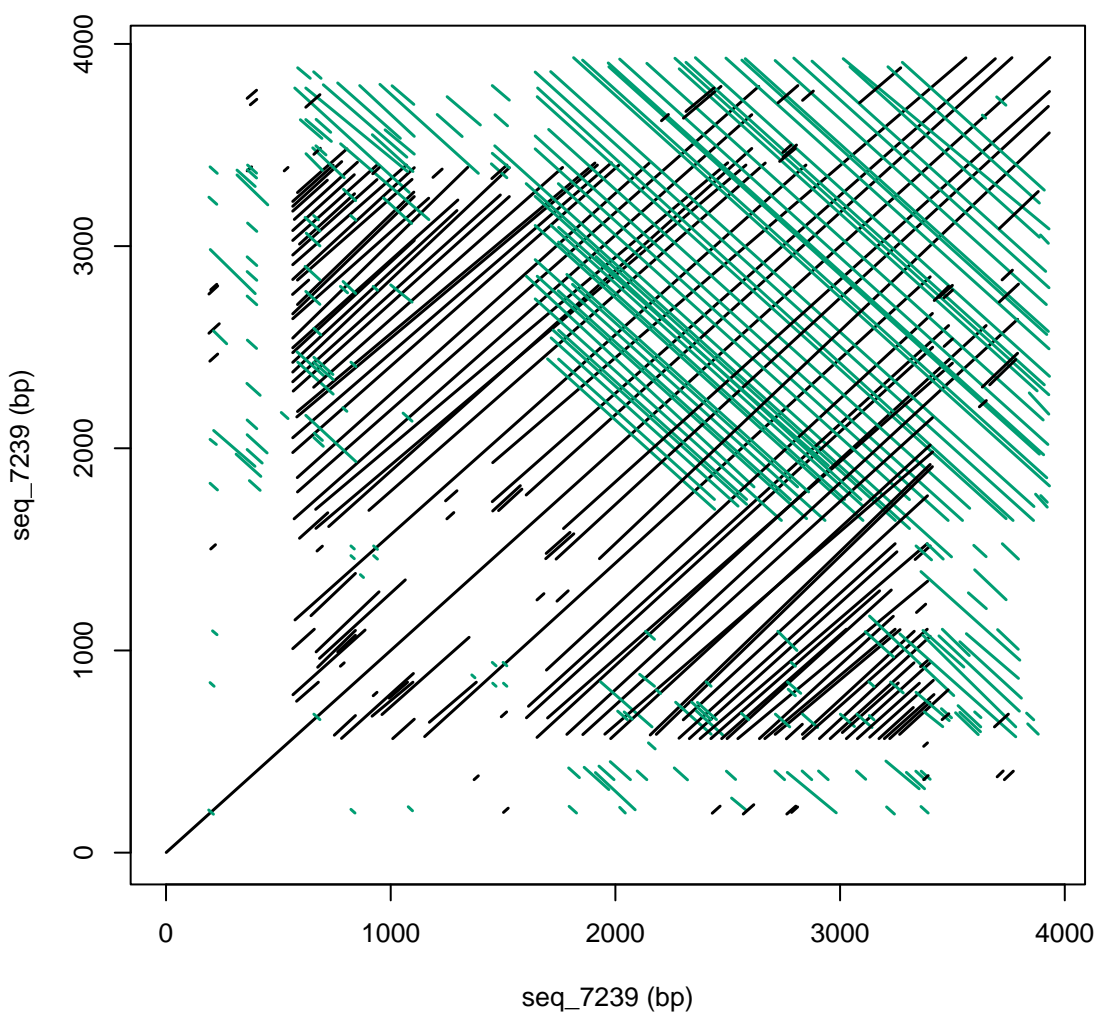
TE: seq_7239
consensus size: 3933bp; fragments: 11535; full length: 188 (≥ 3539.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

