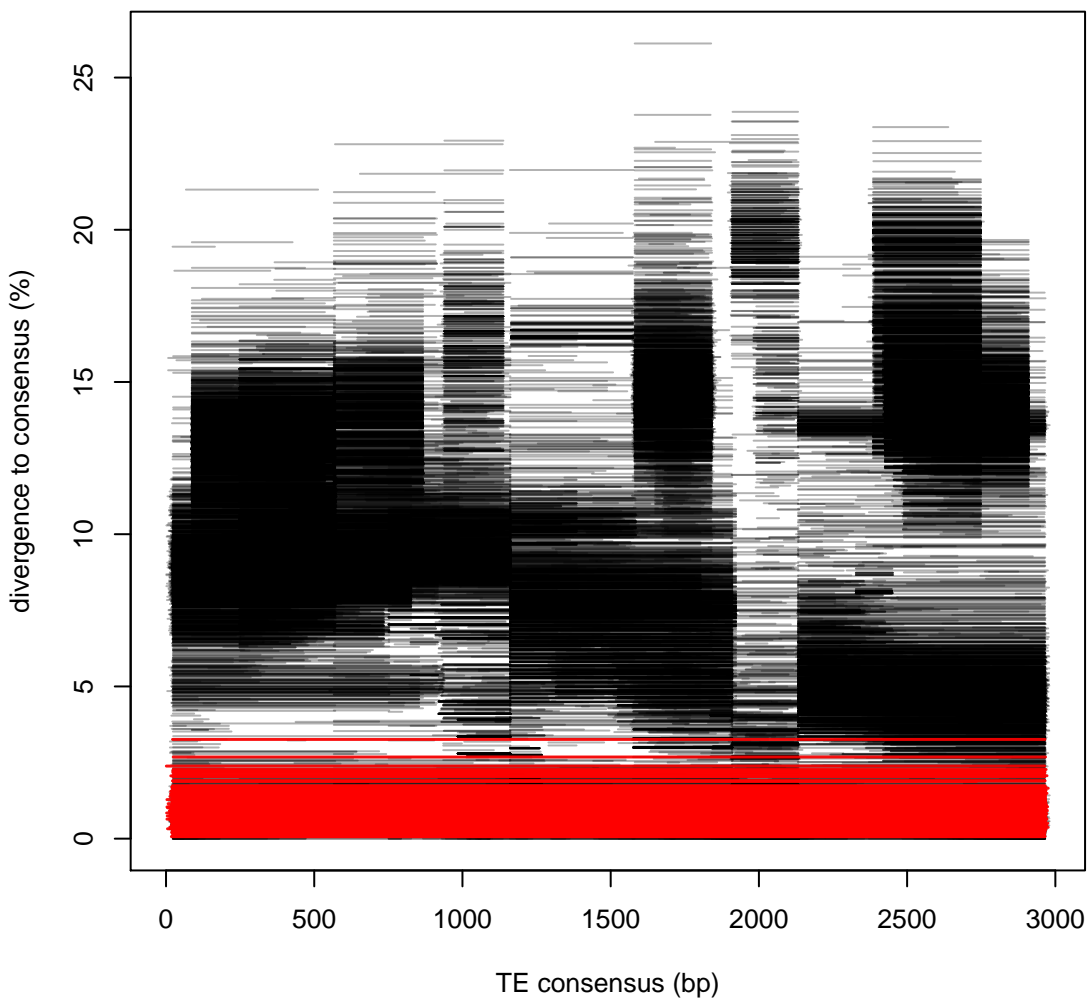
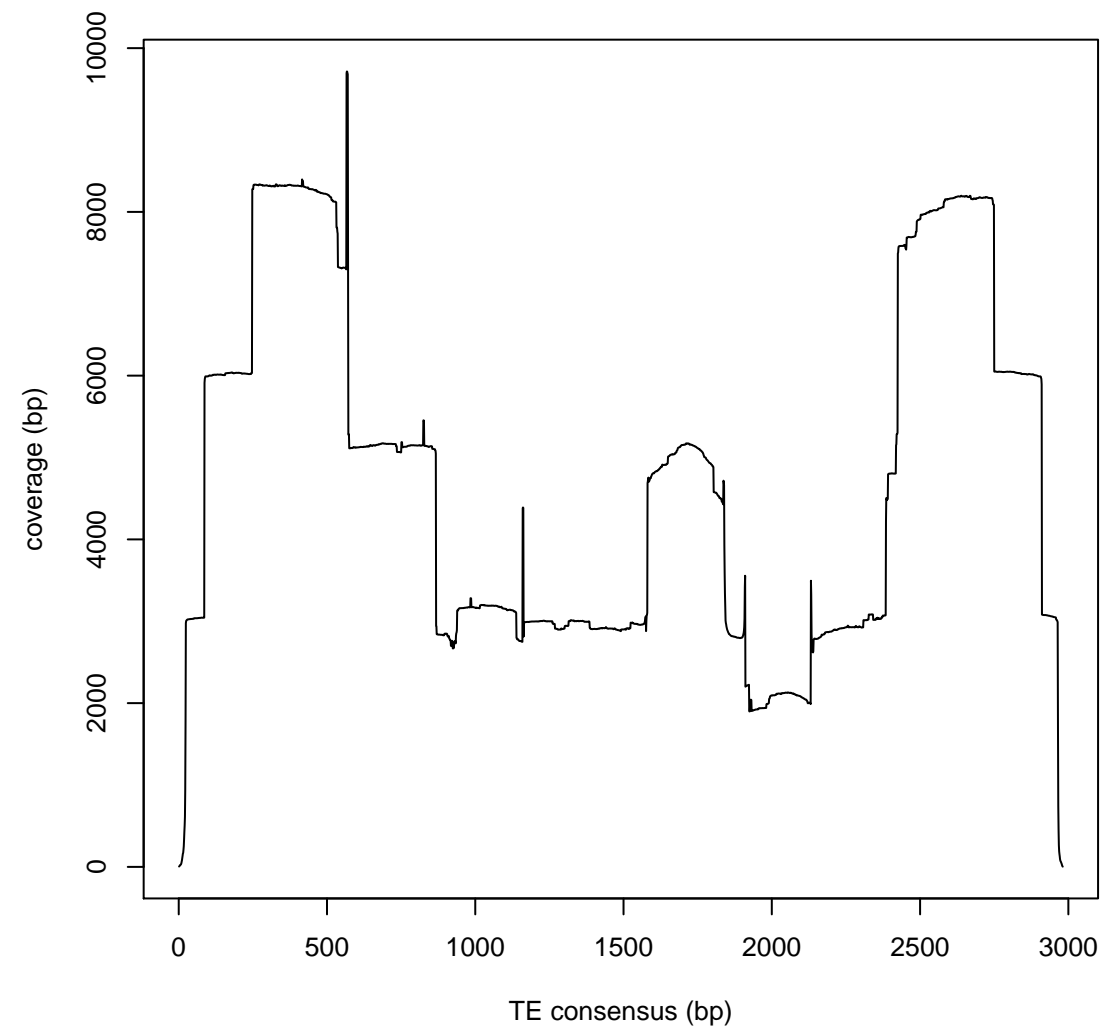


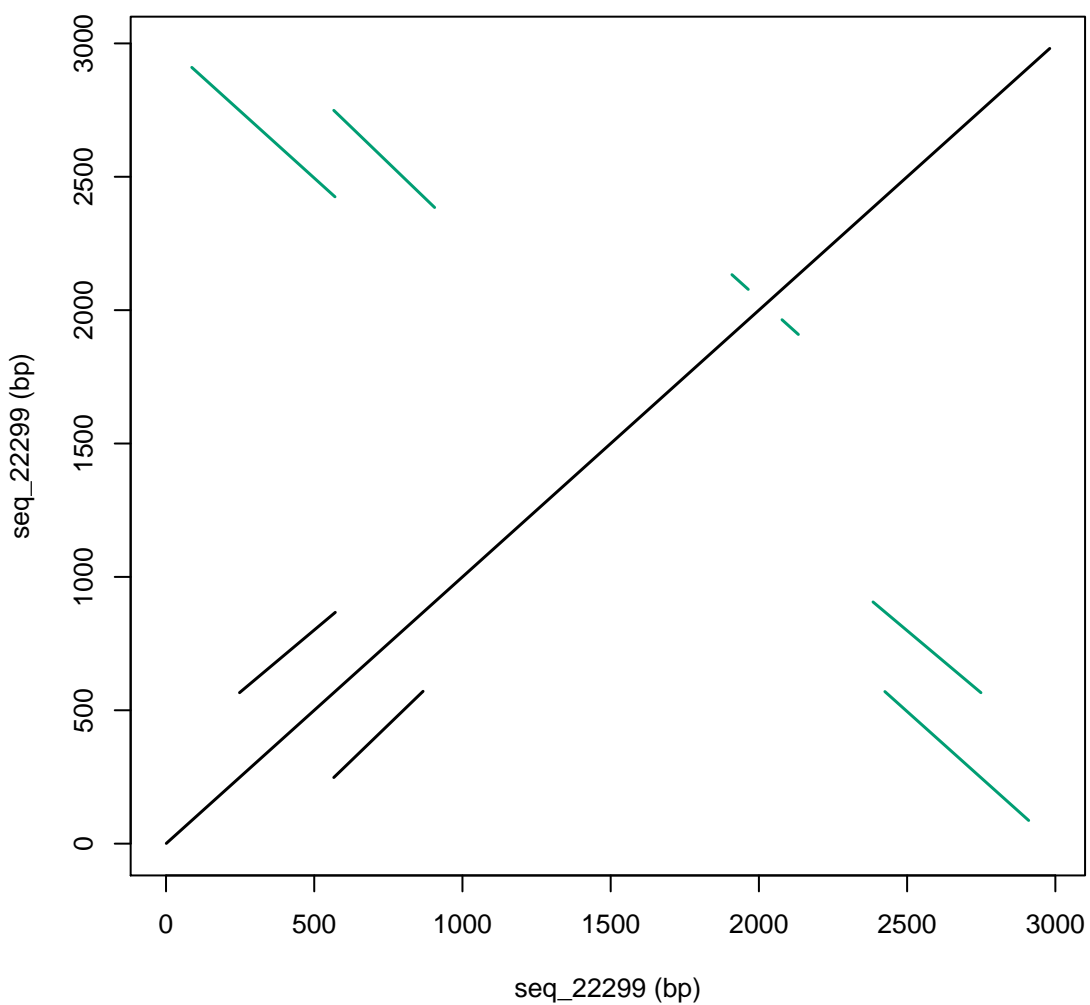
TE: seq_22299
consensus size: 2981bp; fragments: 27872; full length: 454 (≥ 2682.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

