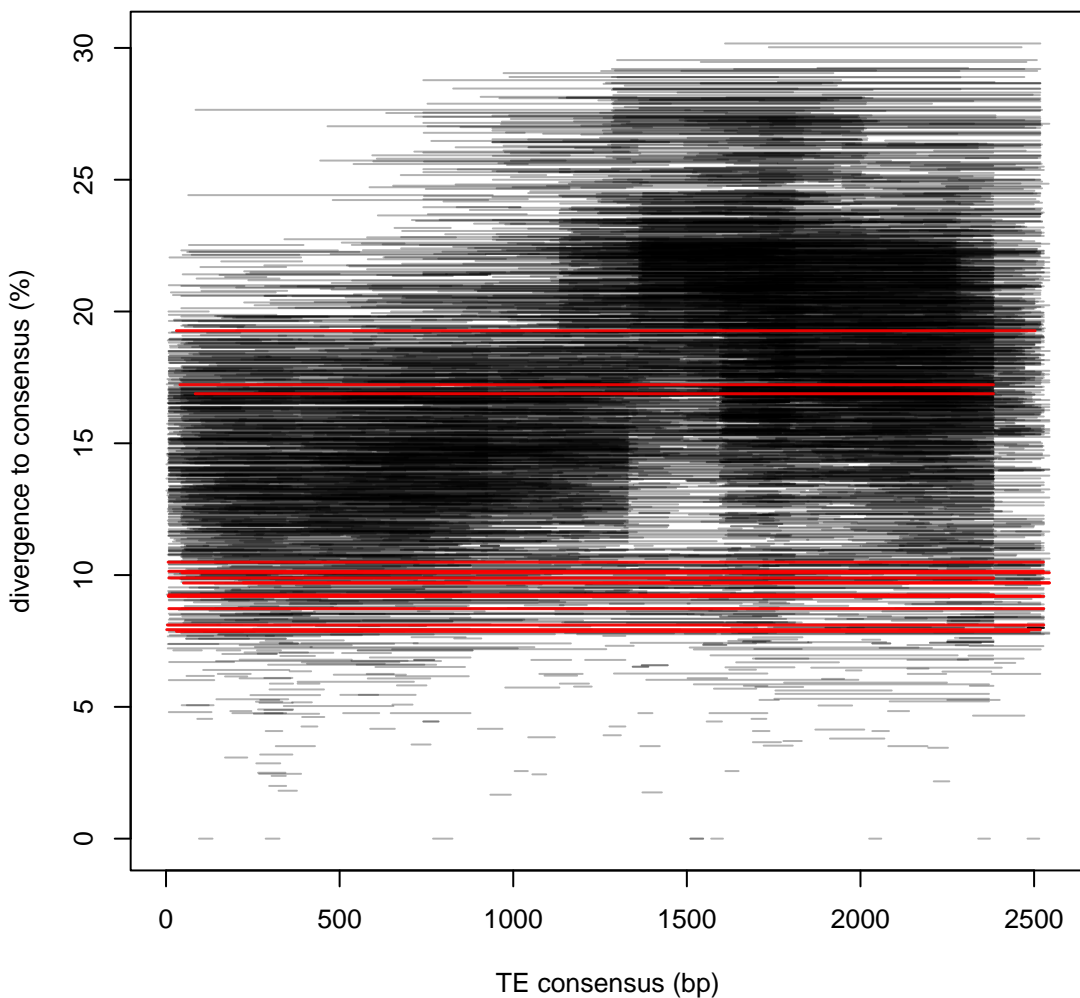
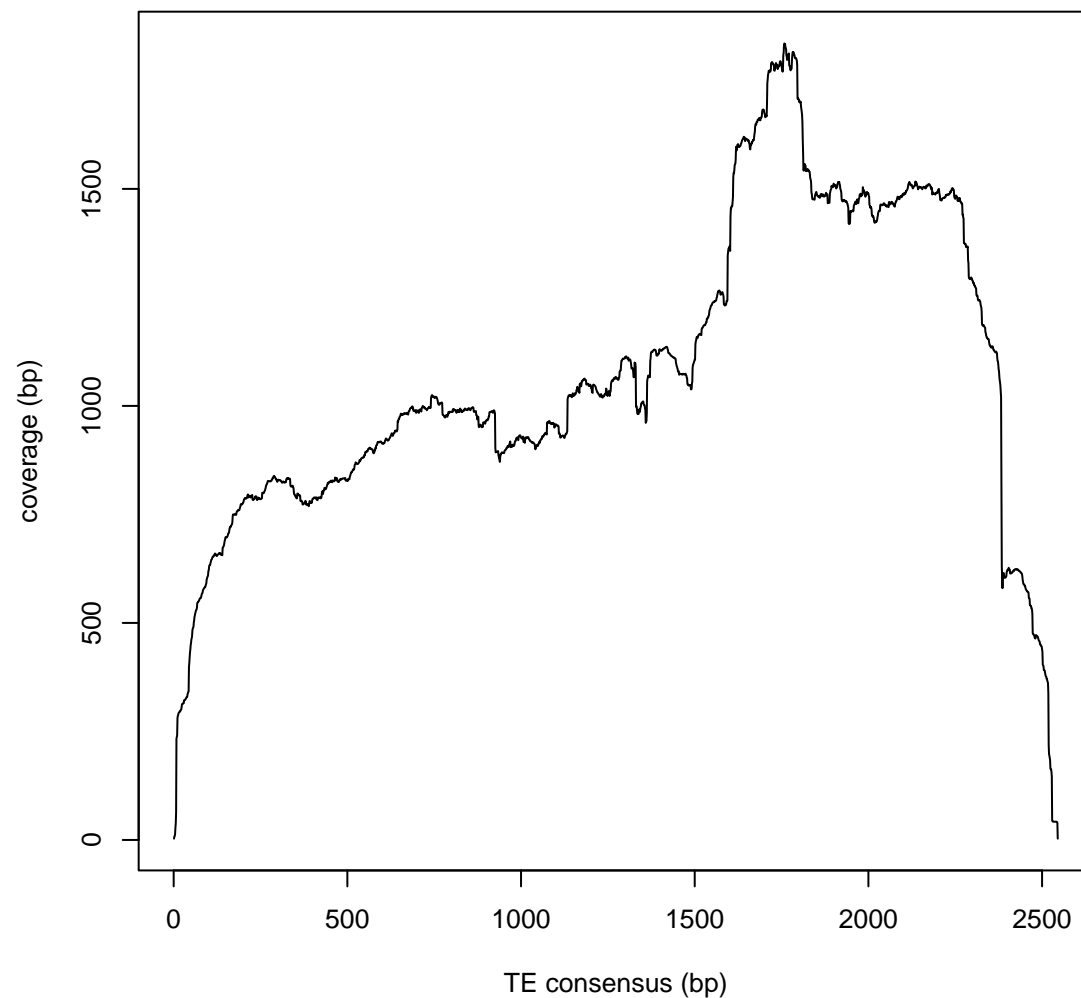


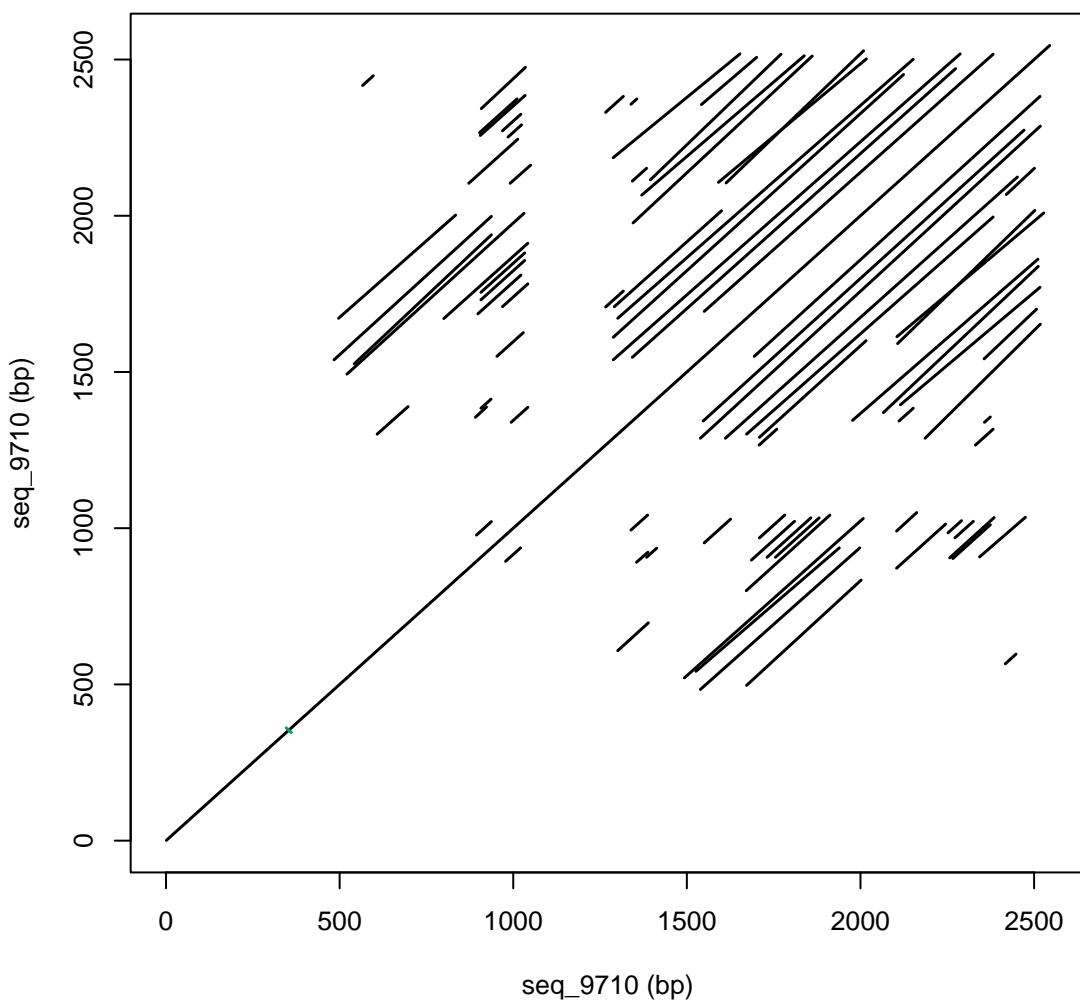
TE: seq_9710
consensus size: 2545bp; fragments: 5301; full length: 16 (≥ 2290.5 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

