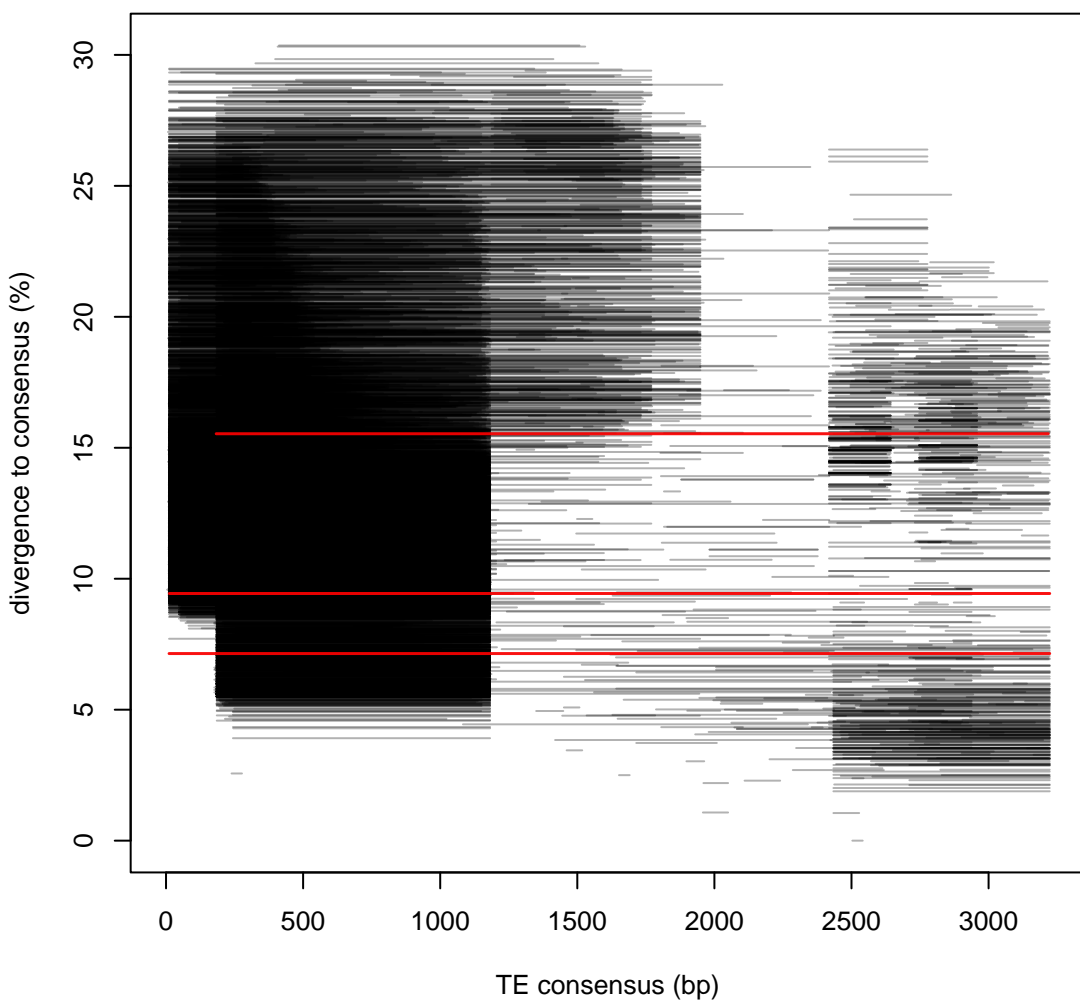
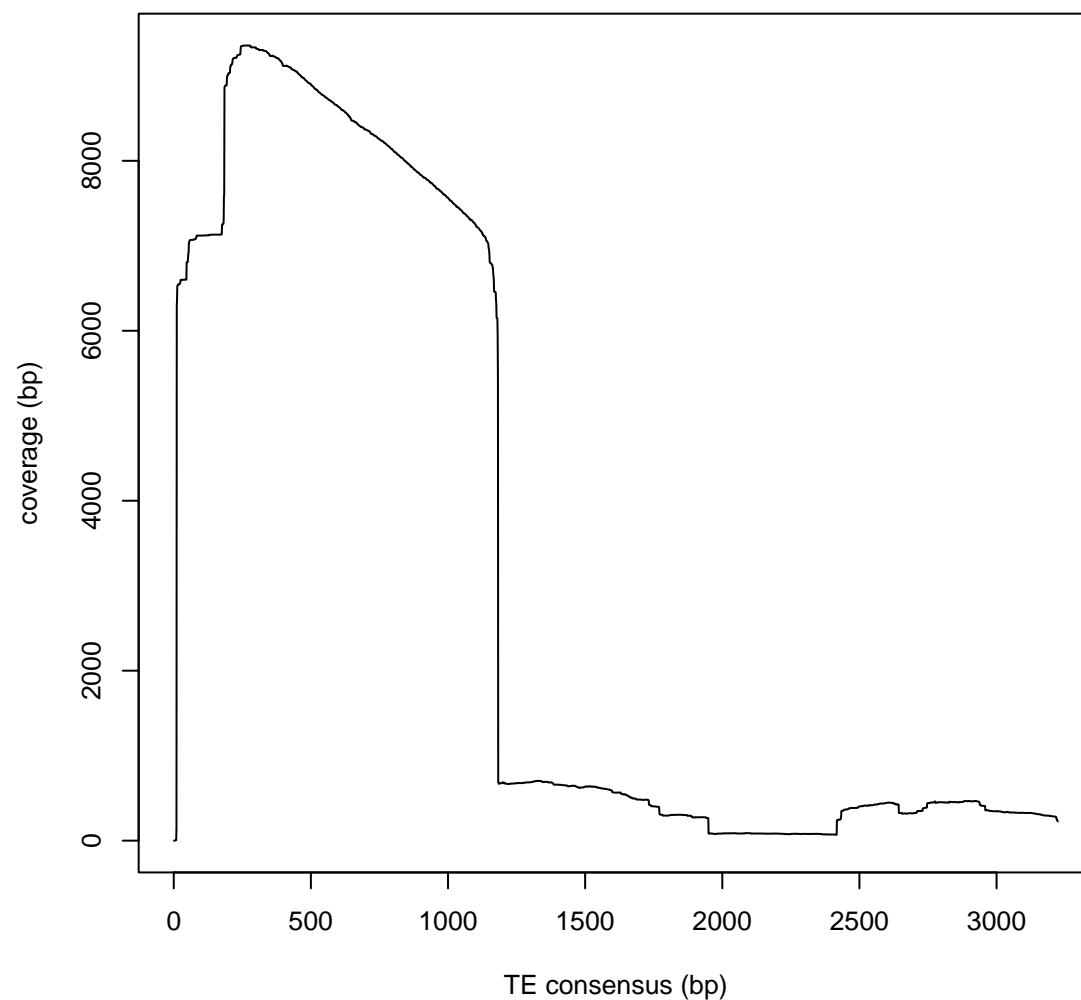


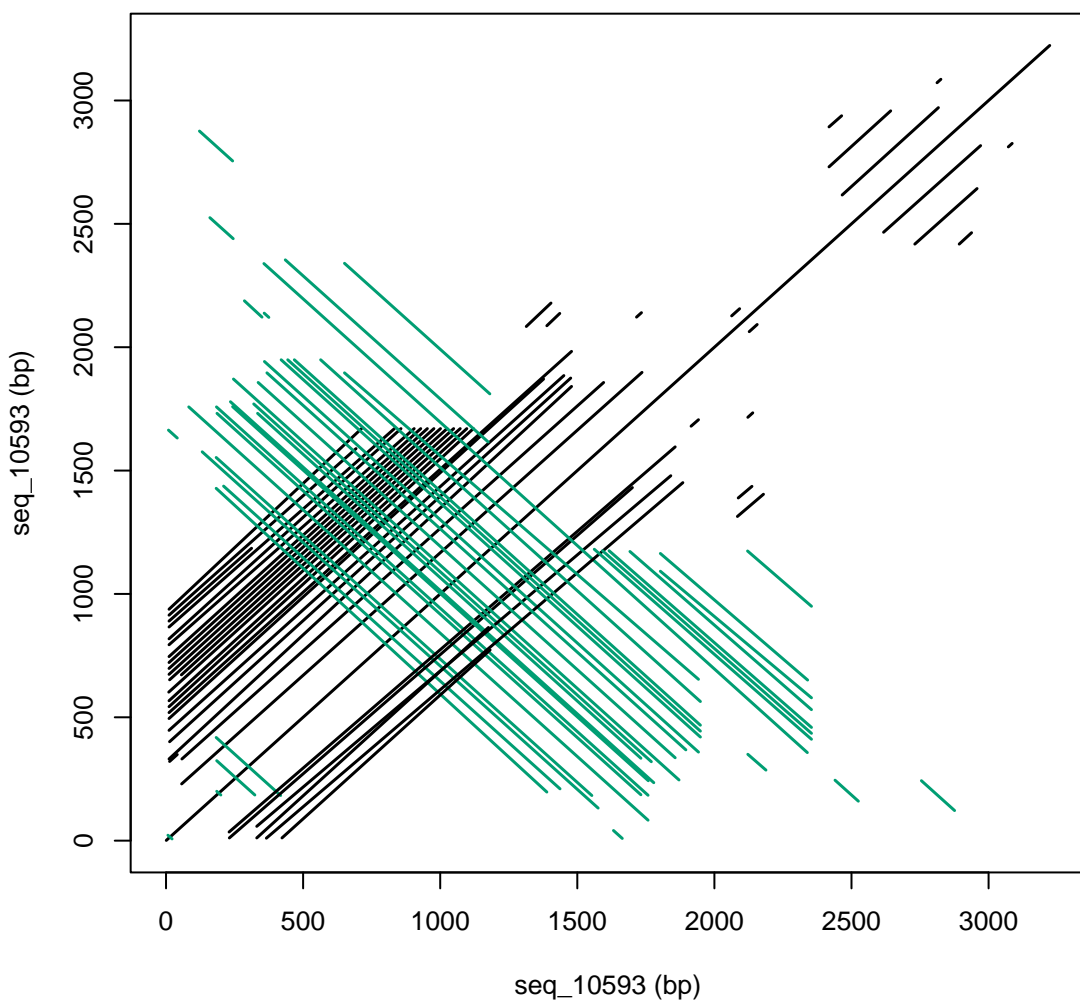
TE: seq_10593
consensus size: 3223bp; fragments: 10358; full length: 3 (>=2900.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

