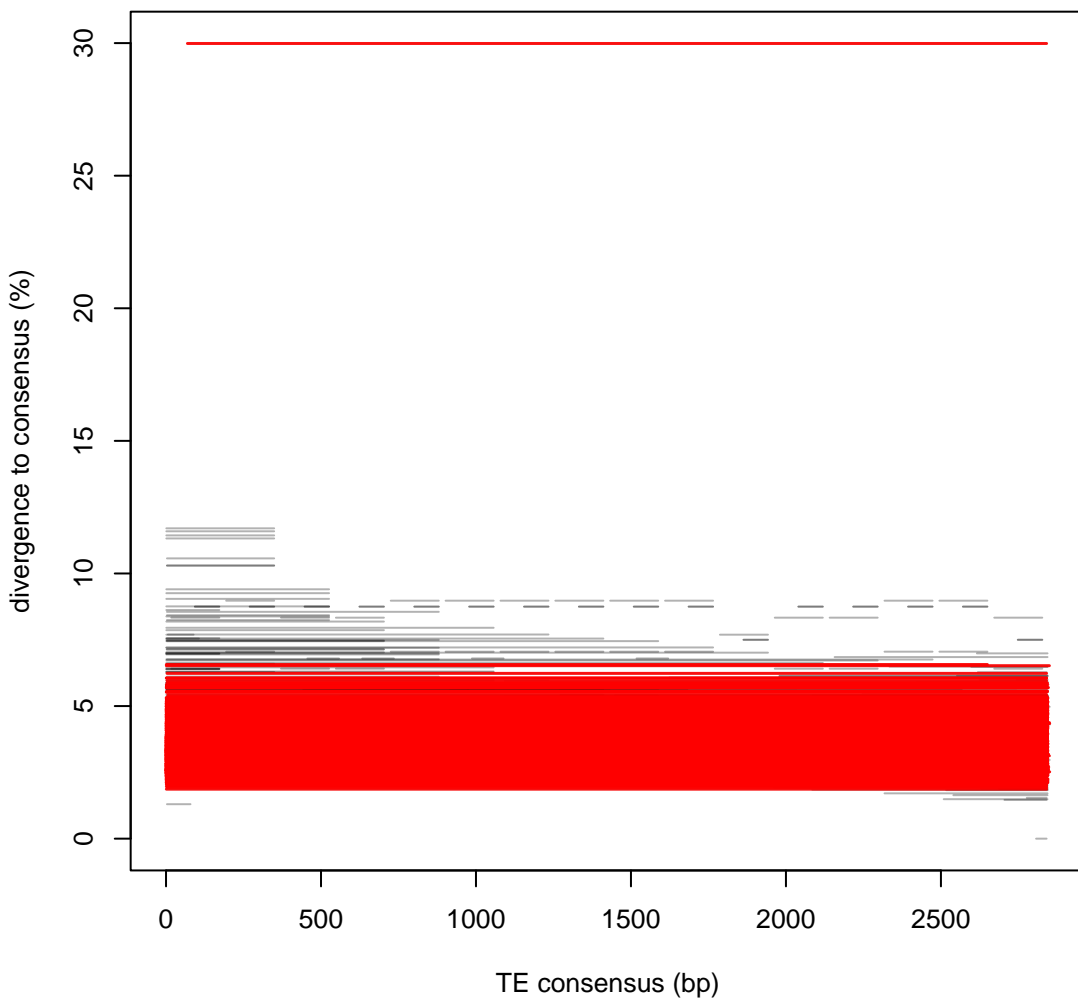
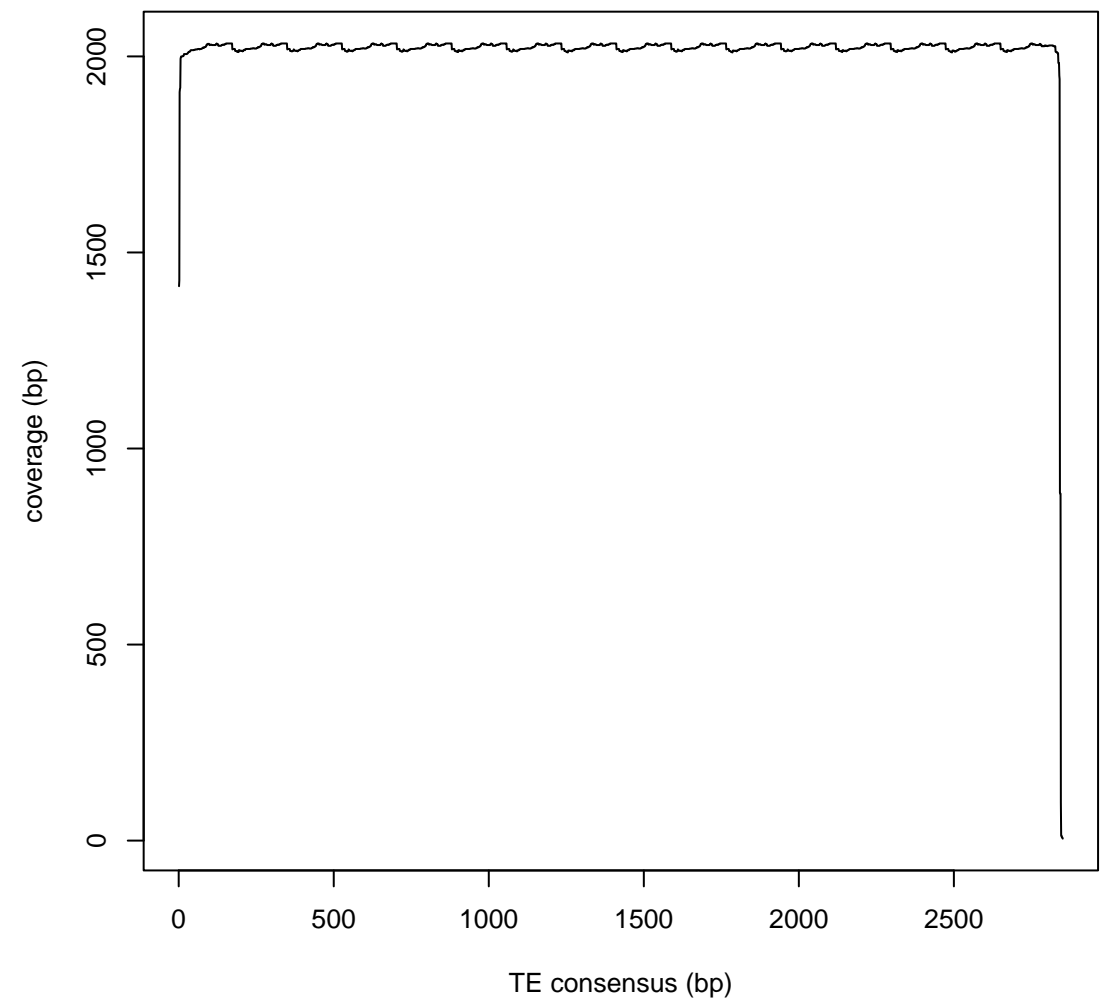


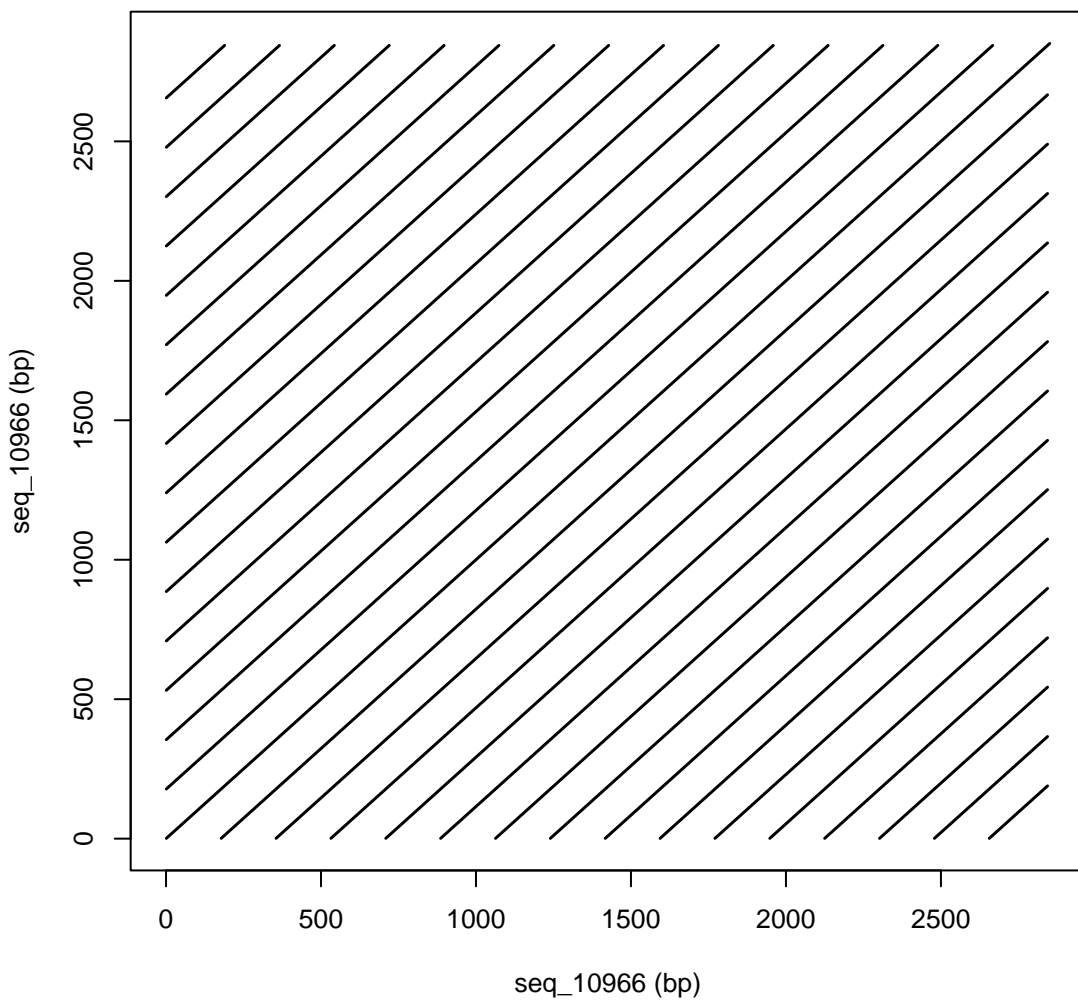
TE: seq_10966
consensus size: 2851bp; fragments: 2719; full length: 1562 (≥ 2565.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

