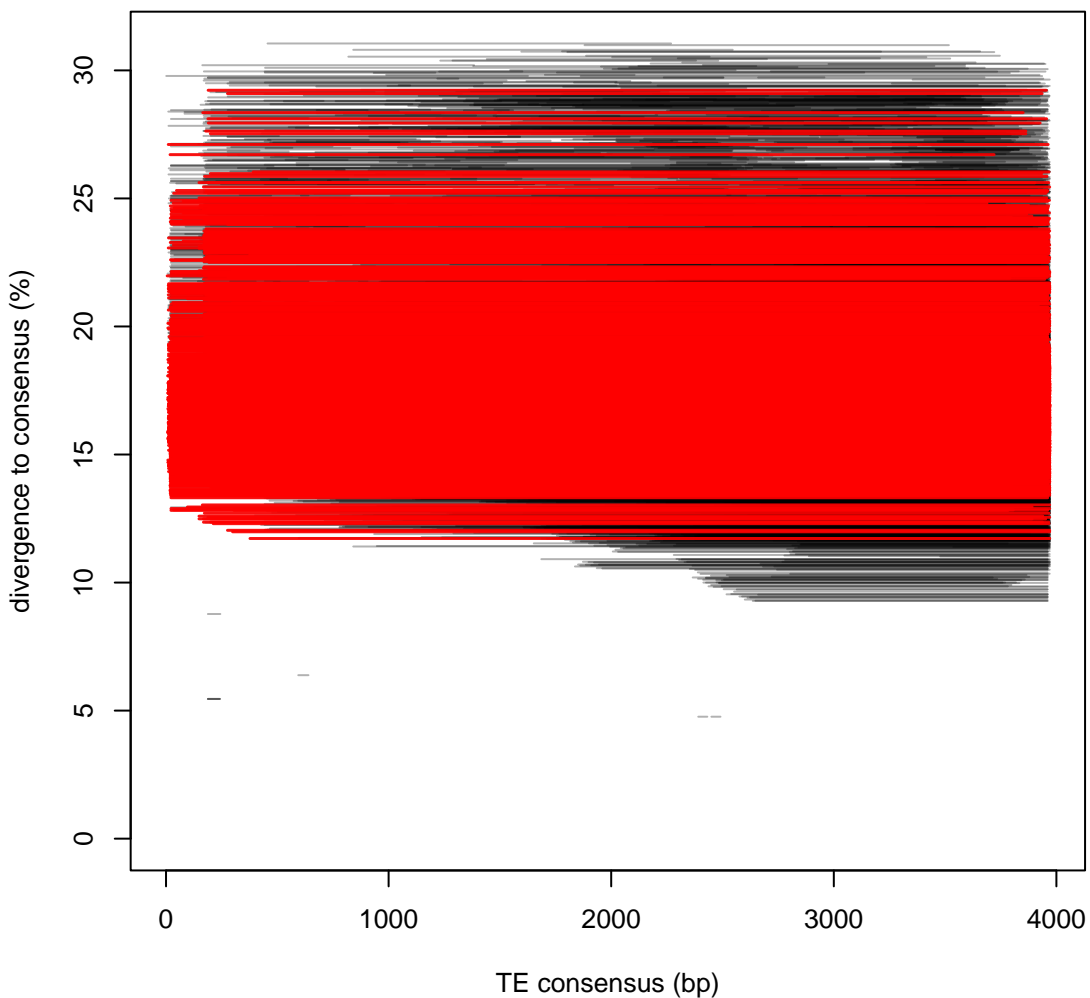
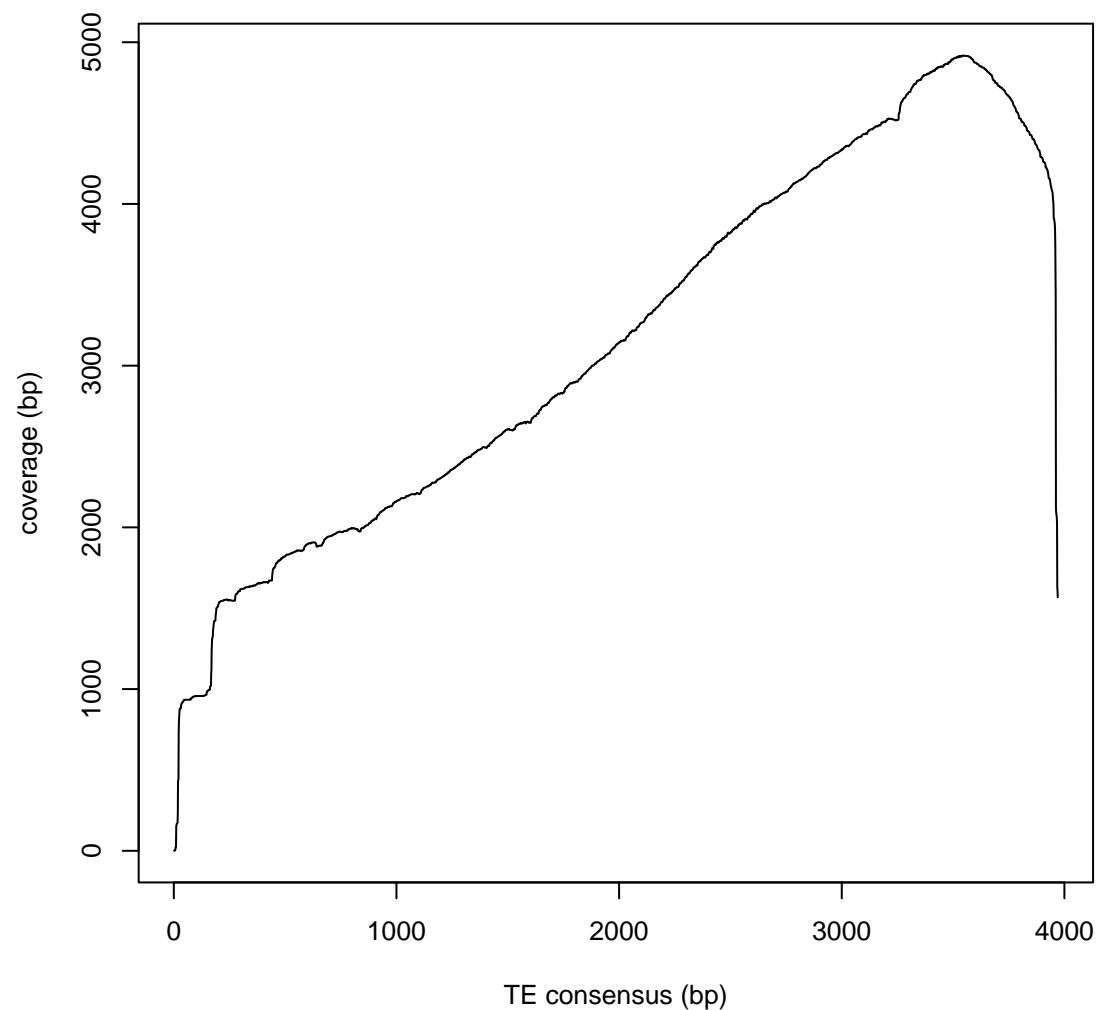


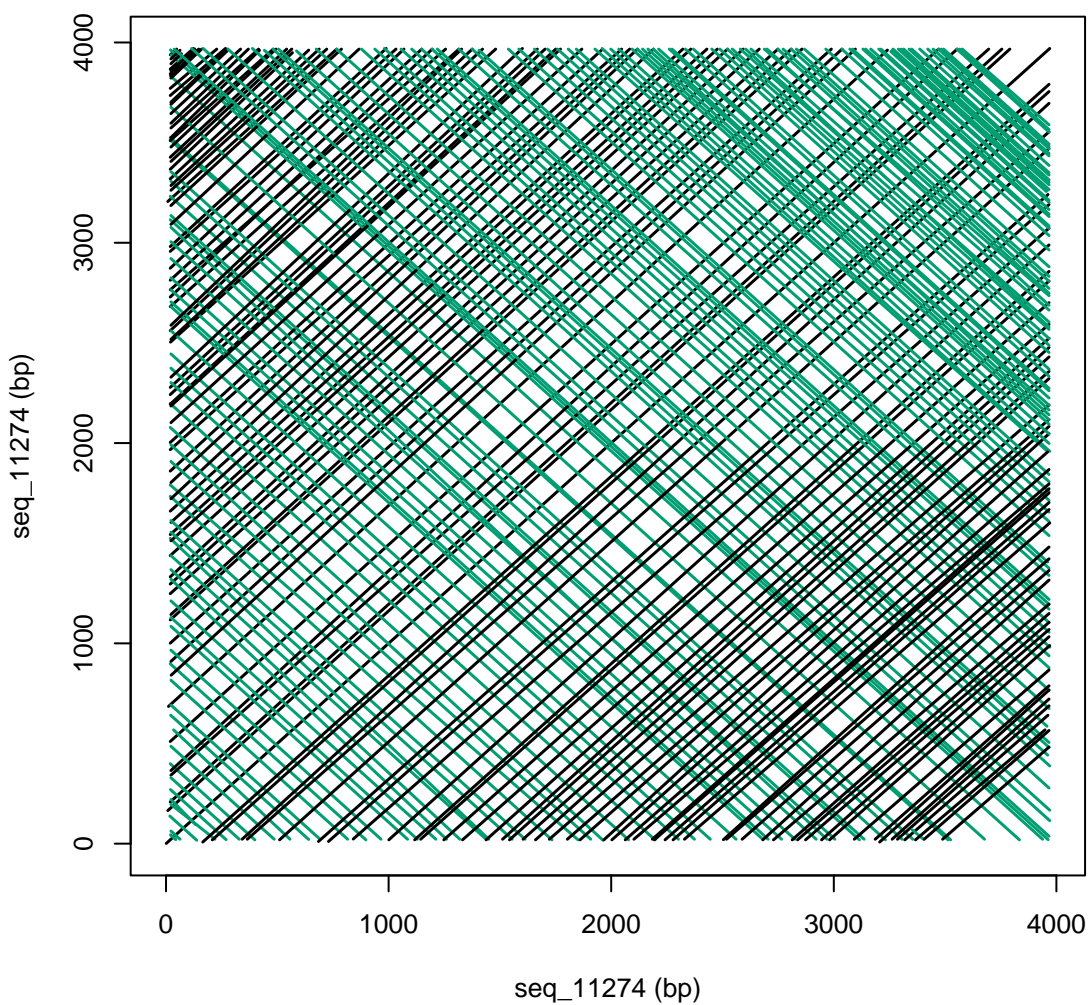
TE: seq_11274
consensus size: 3970bp; fragments: 5795; full length: 1160 (≥ 3573 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

