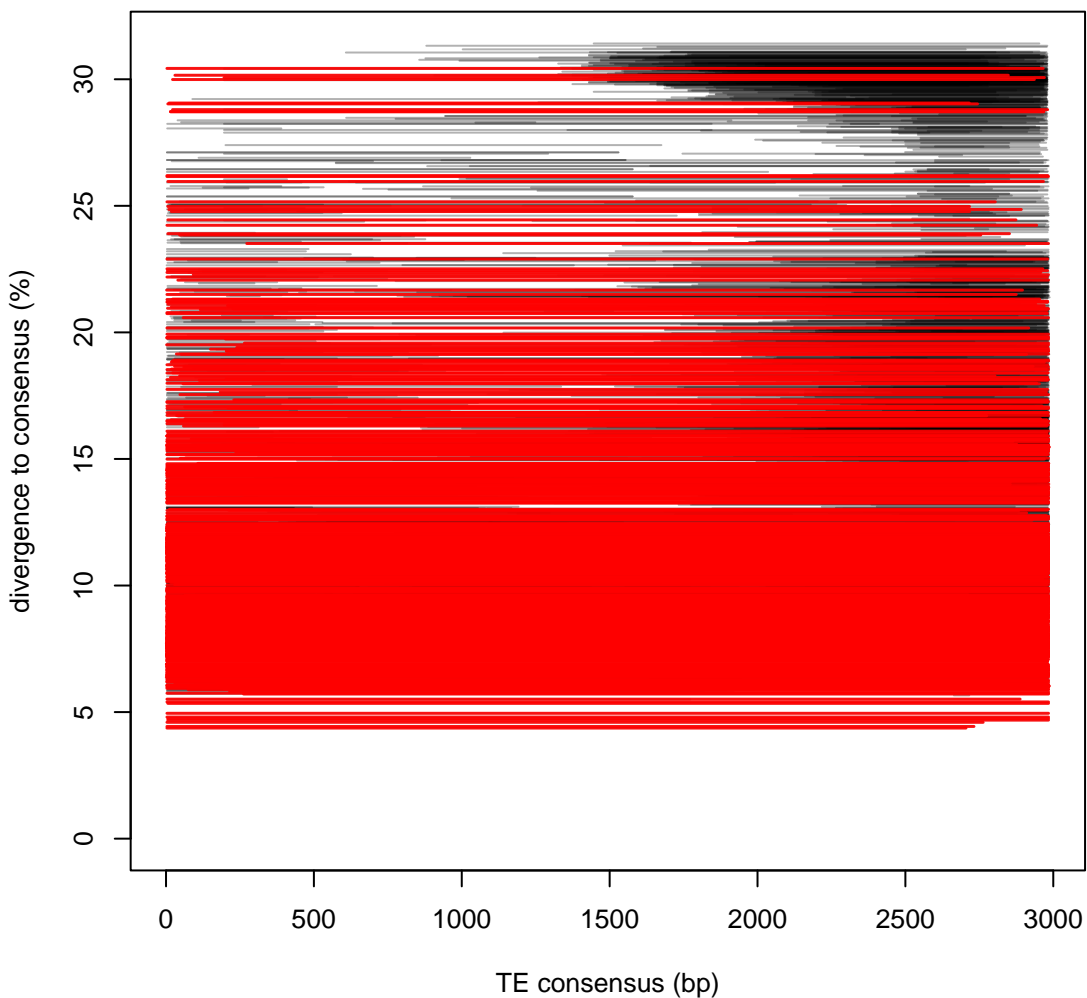
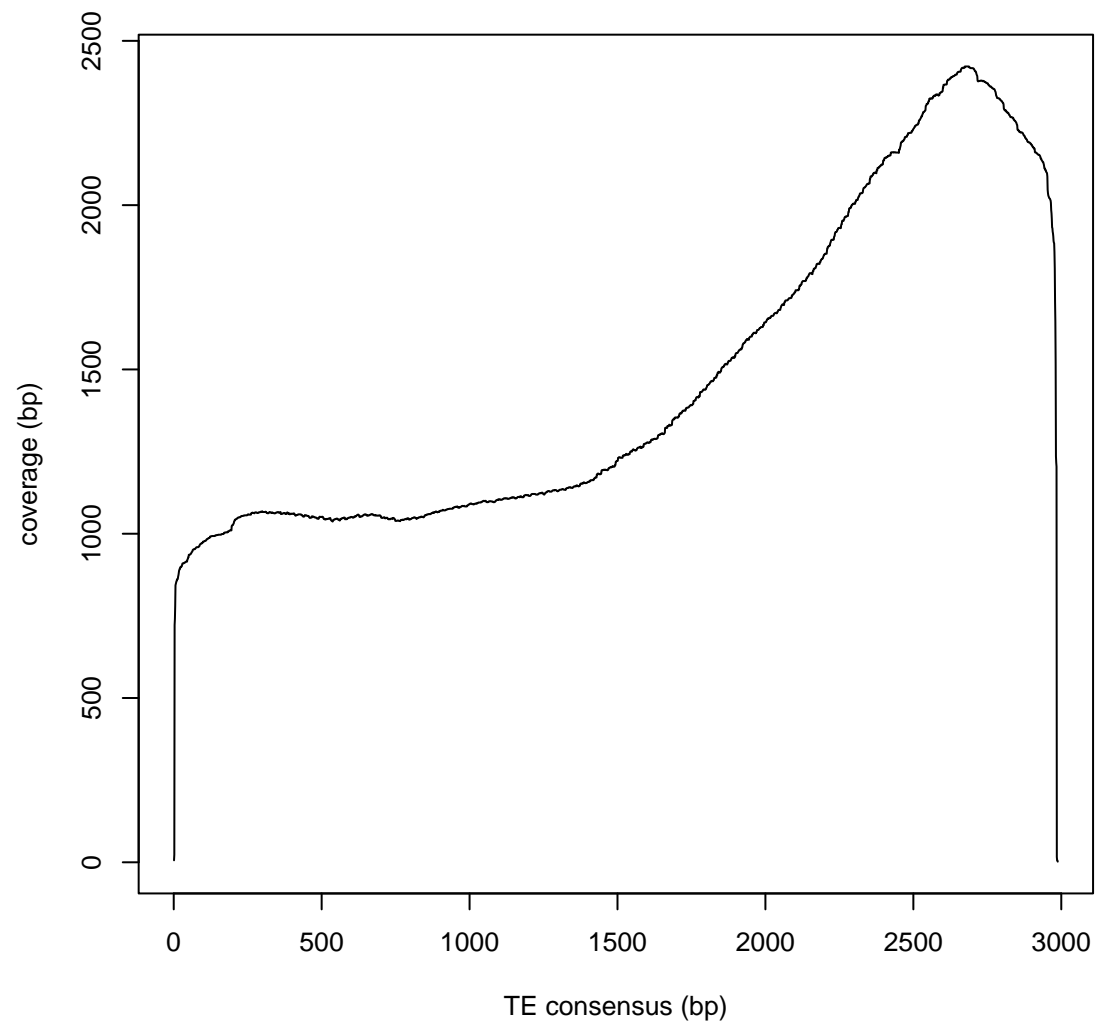


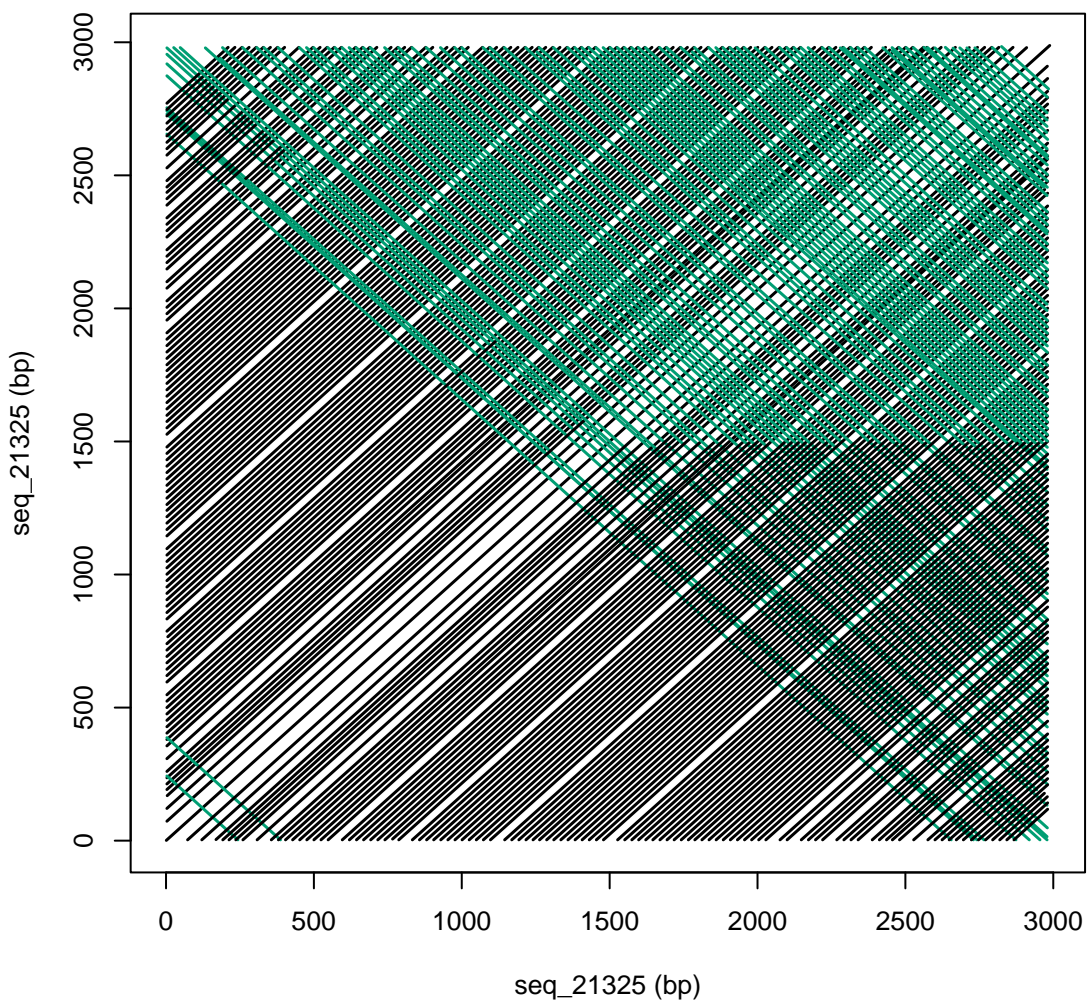
TE: seq_21325
consensus size: 2988bp; fragments: 2846; full length: 656 (≥ 2689.2 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

