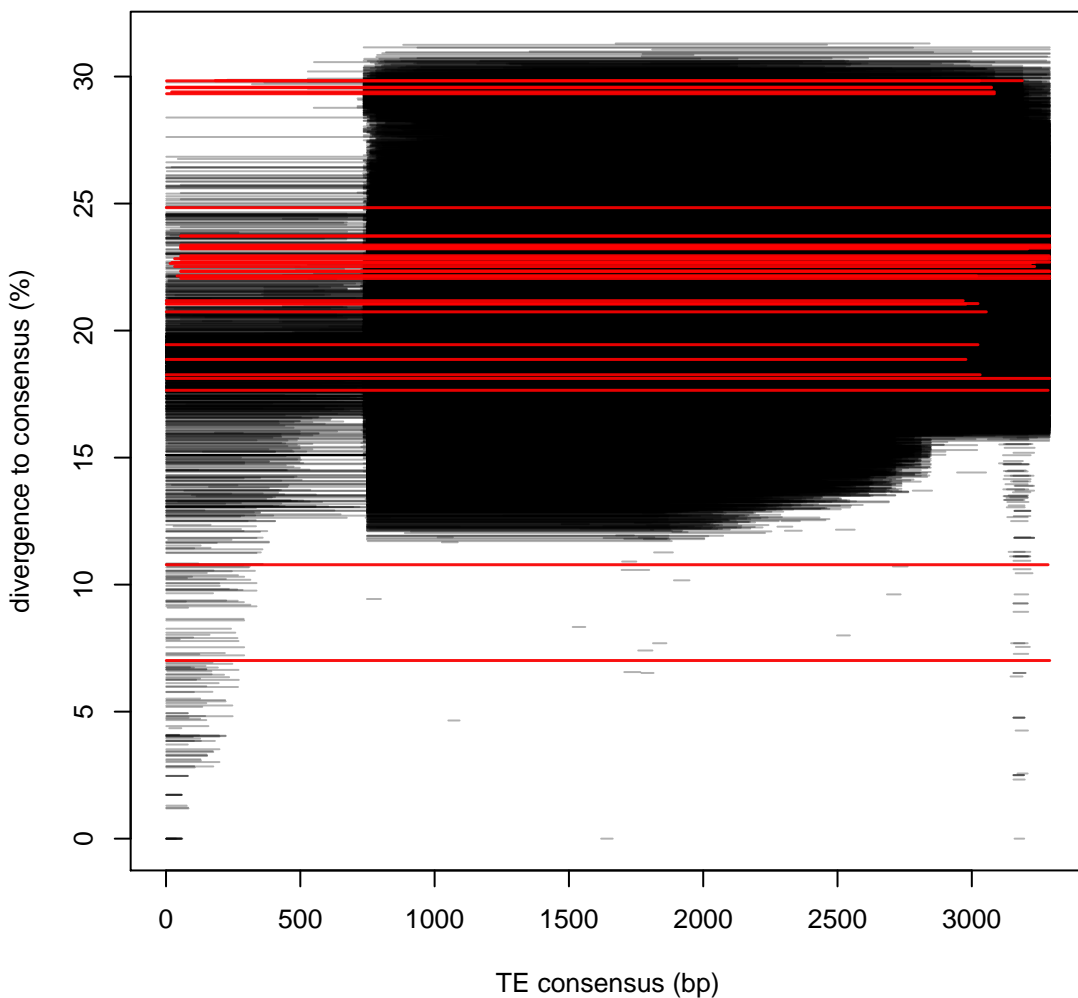
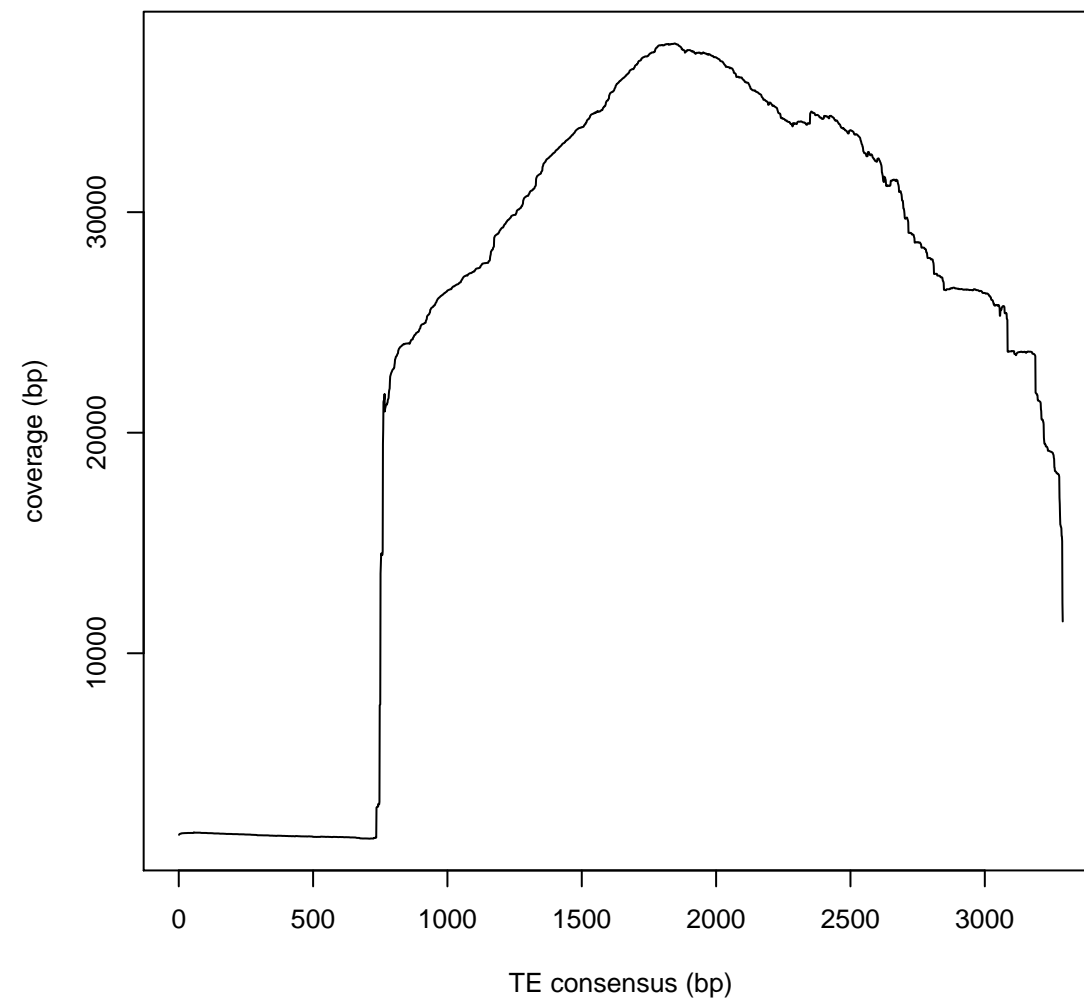


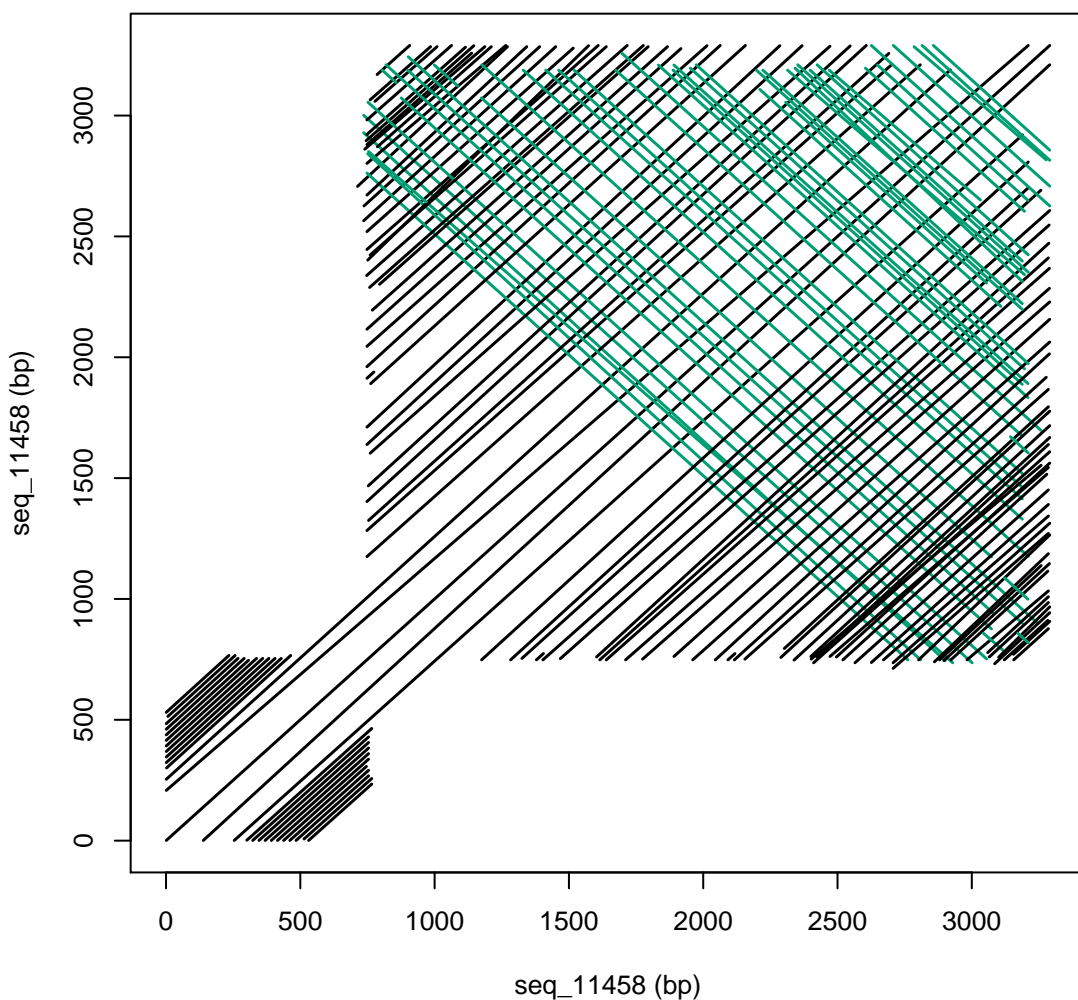
TE: seq_11458
consensus size: 3290bp; fragments: 46454; full length: 39 (>=2961bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

