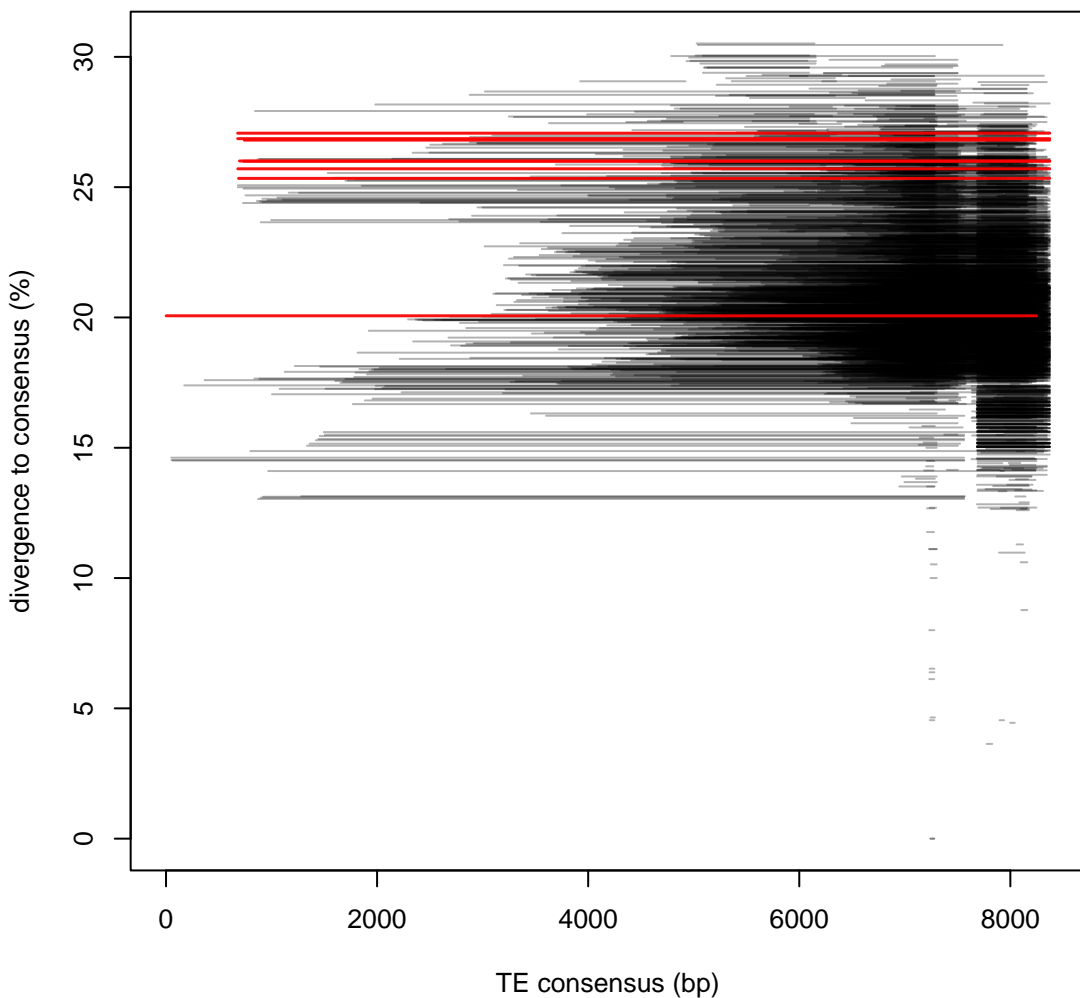
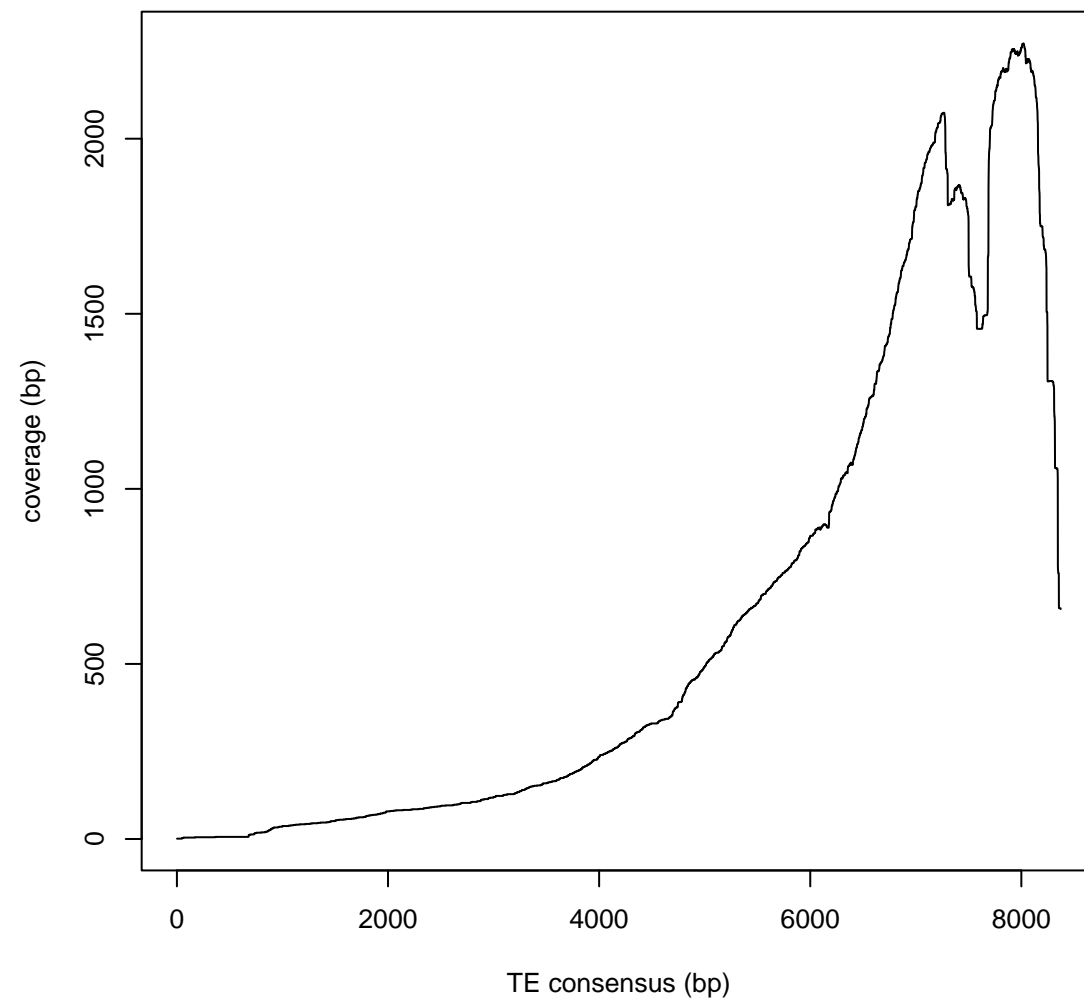


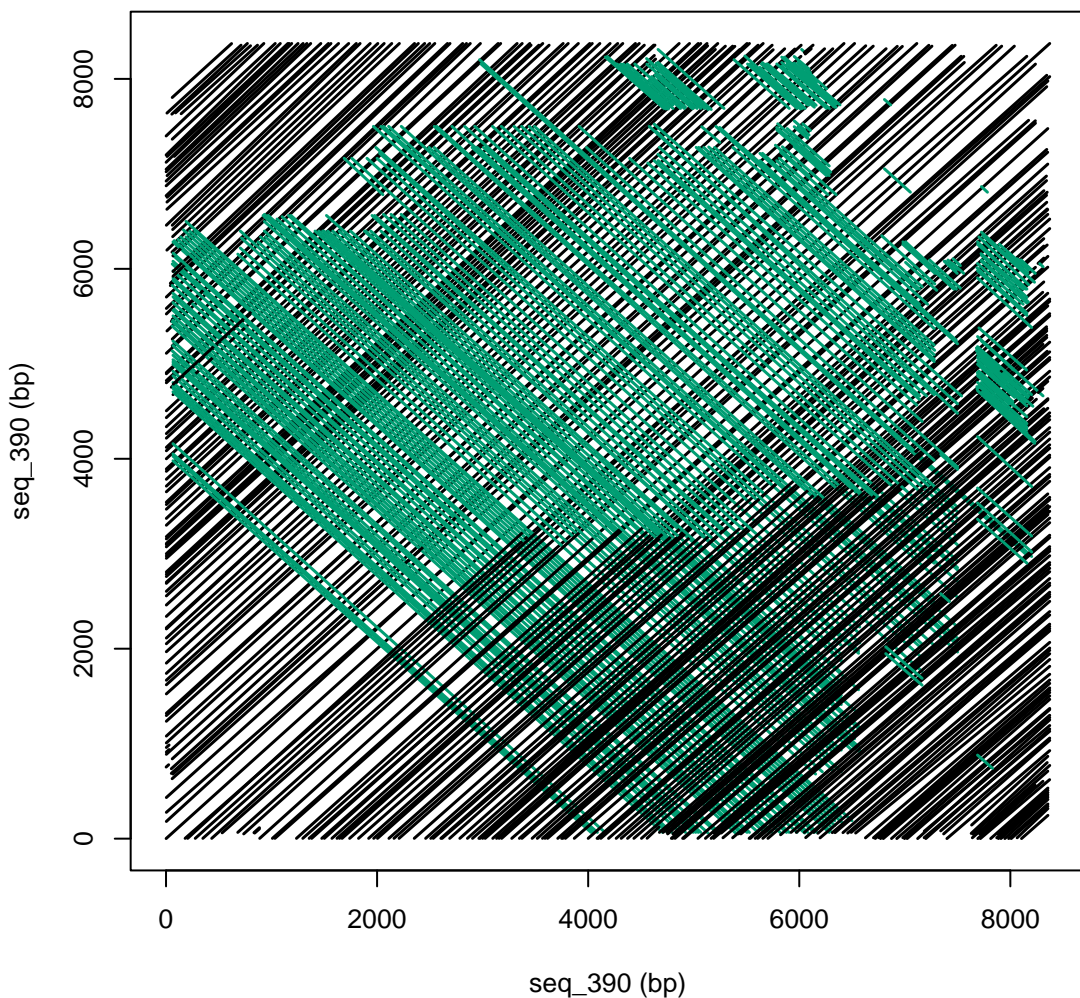
TE: seq_390
consensus size: 8373bp; fragments: 3184; full length: 8 (>=7535.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

