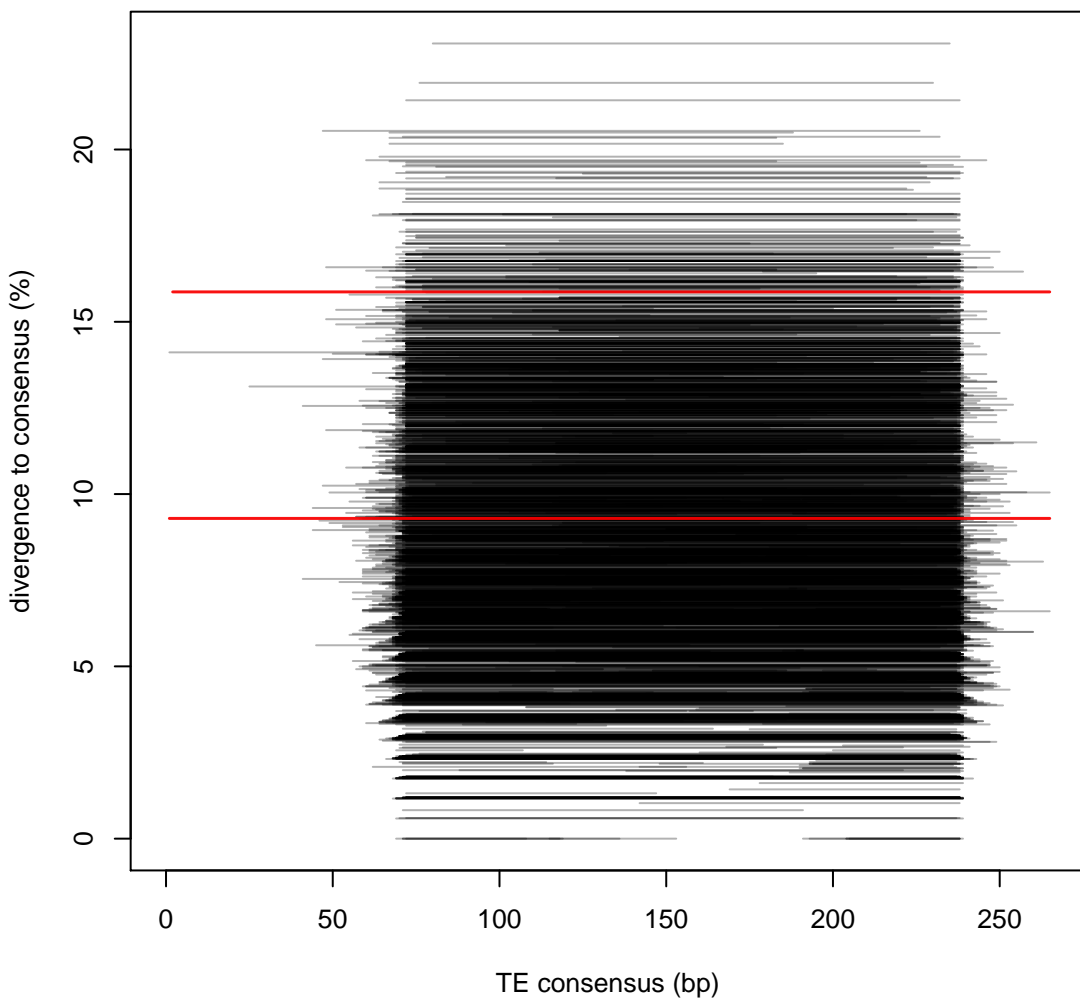
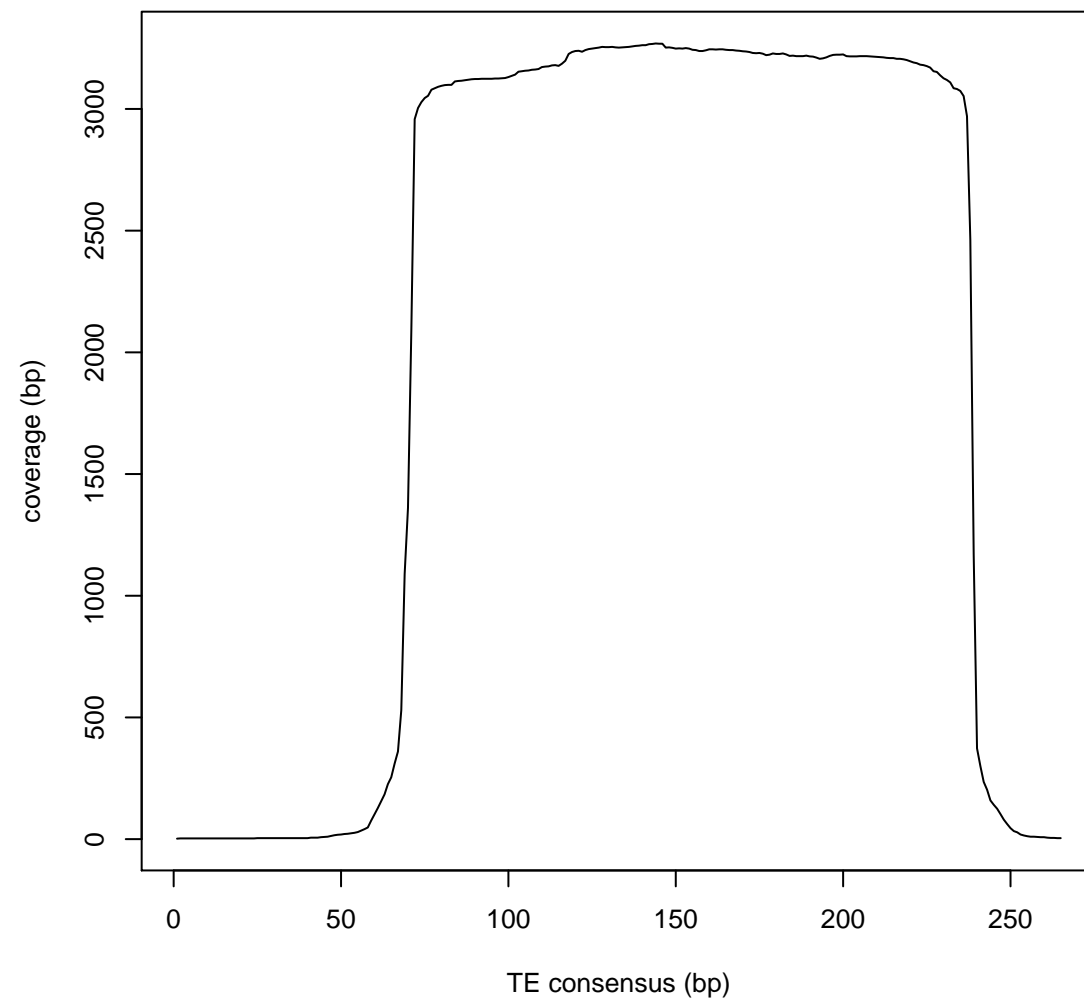


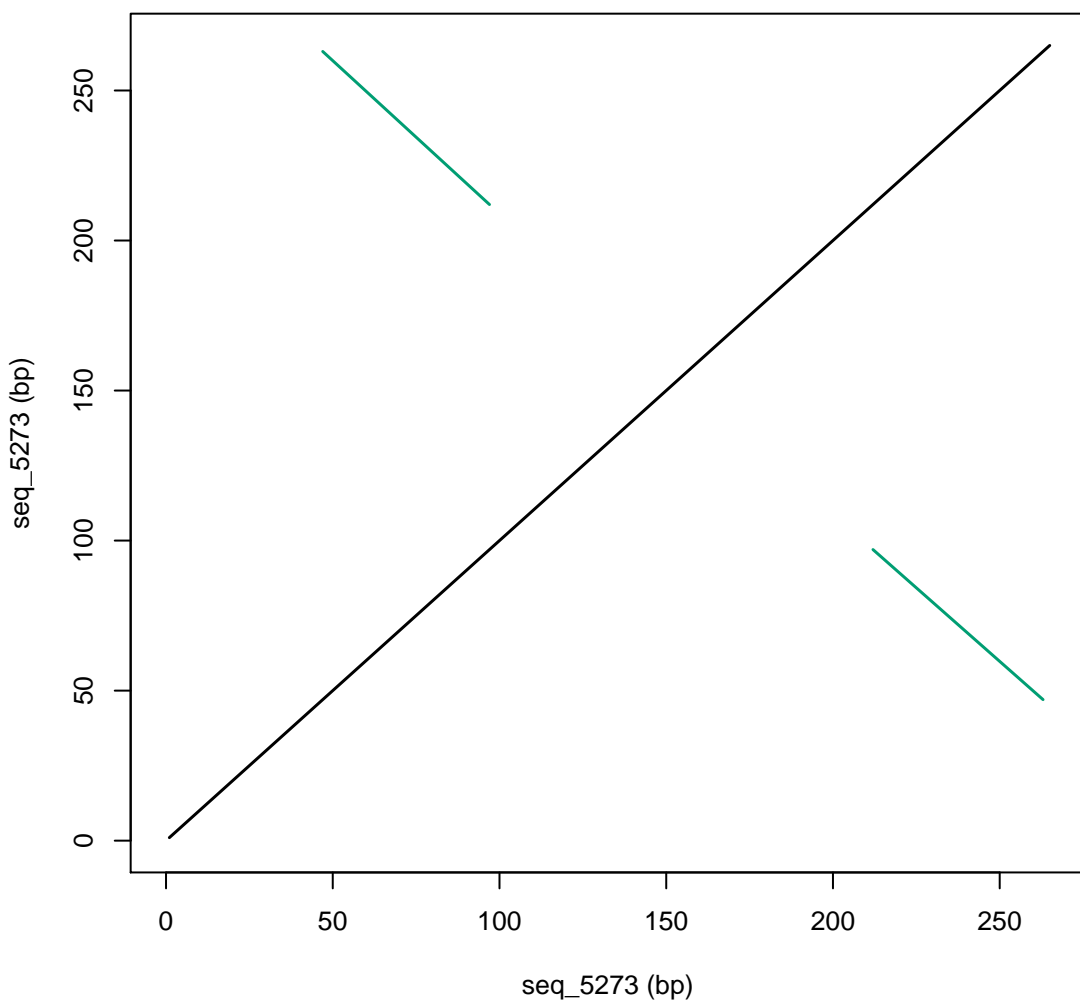
TE: seq_5273
consensus size: 265bp; fragments: 3435; full length: 2 (≥ 238.5 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

