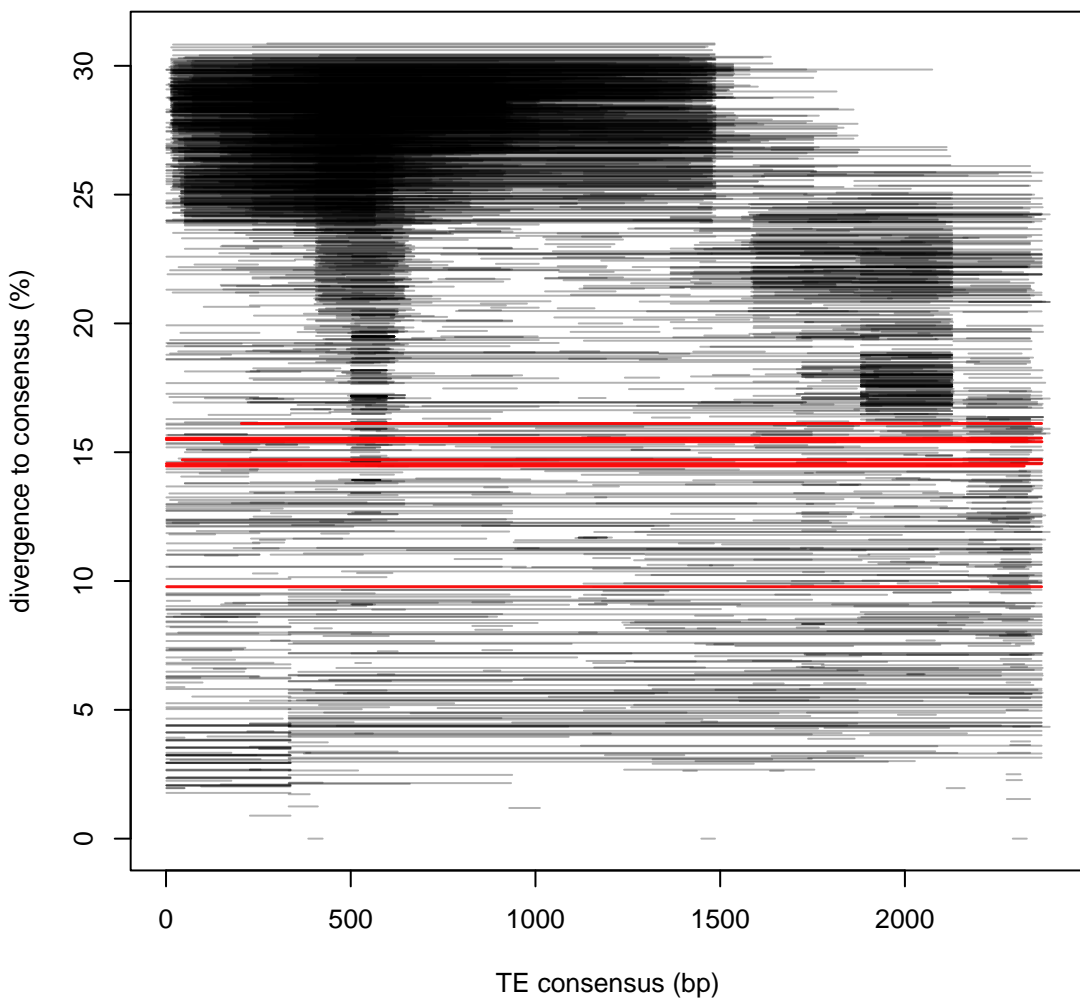
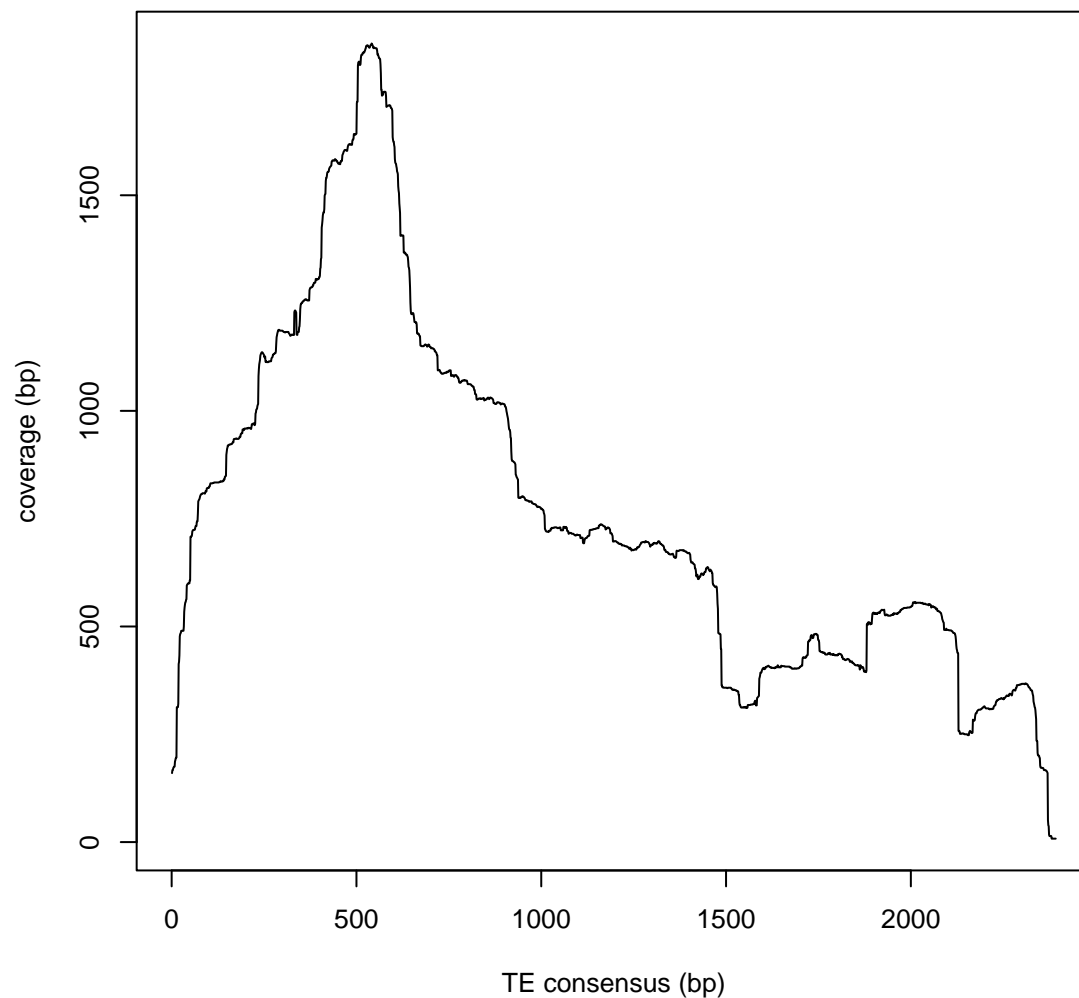


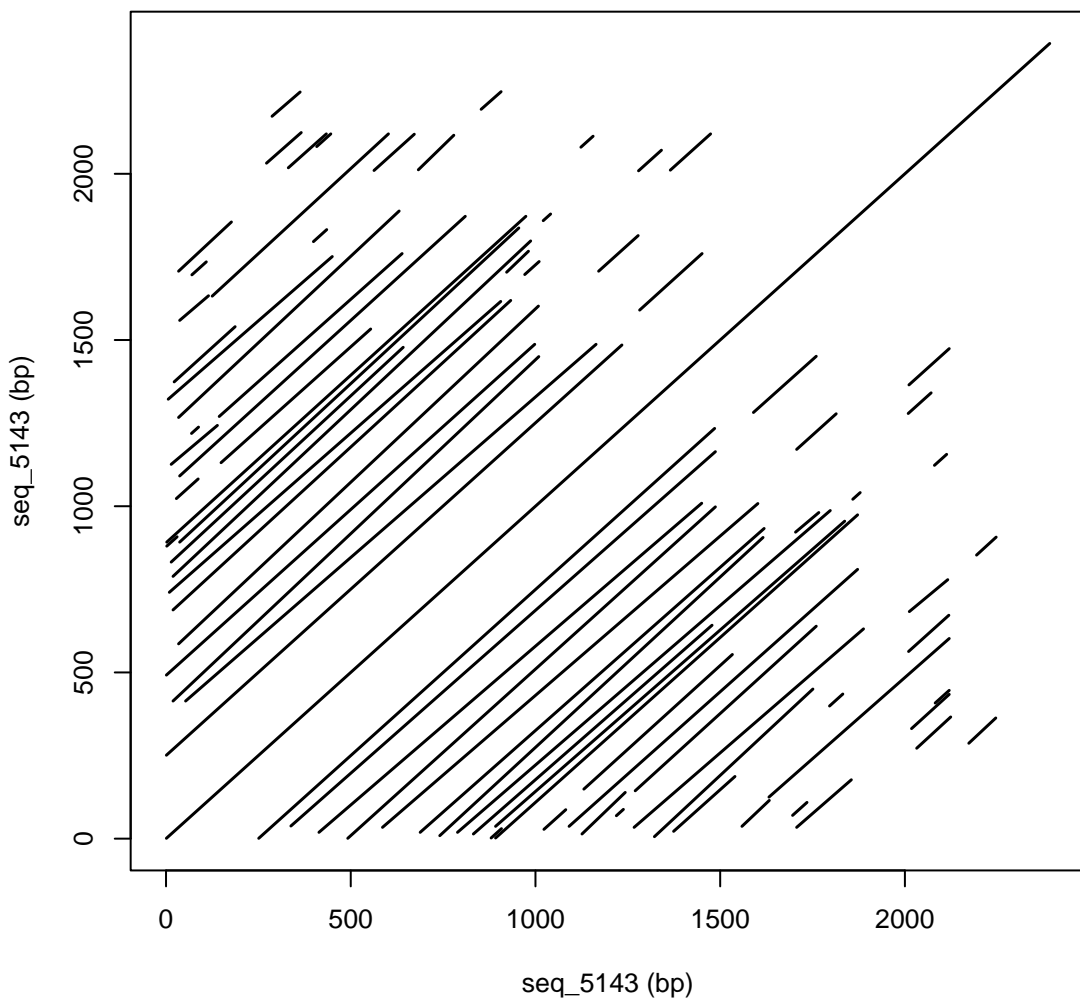
TE: seq_5143
consensus size: 2392bp; fragments: 3131; full length: 8 (≥ 2152.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

