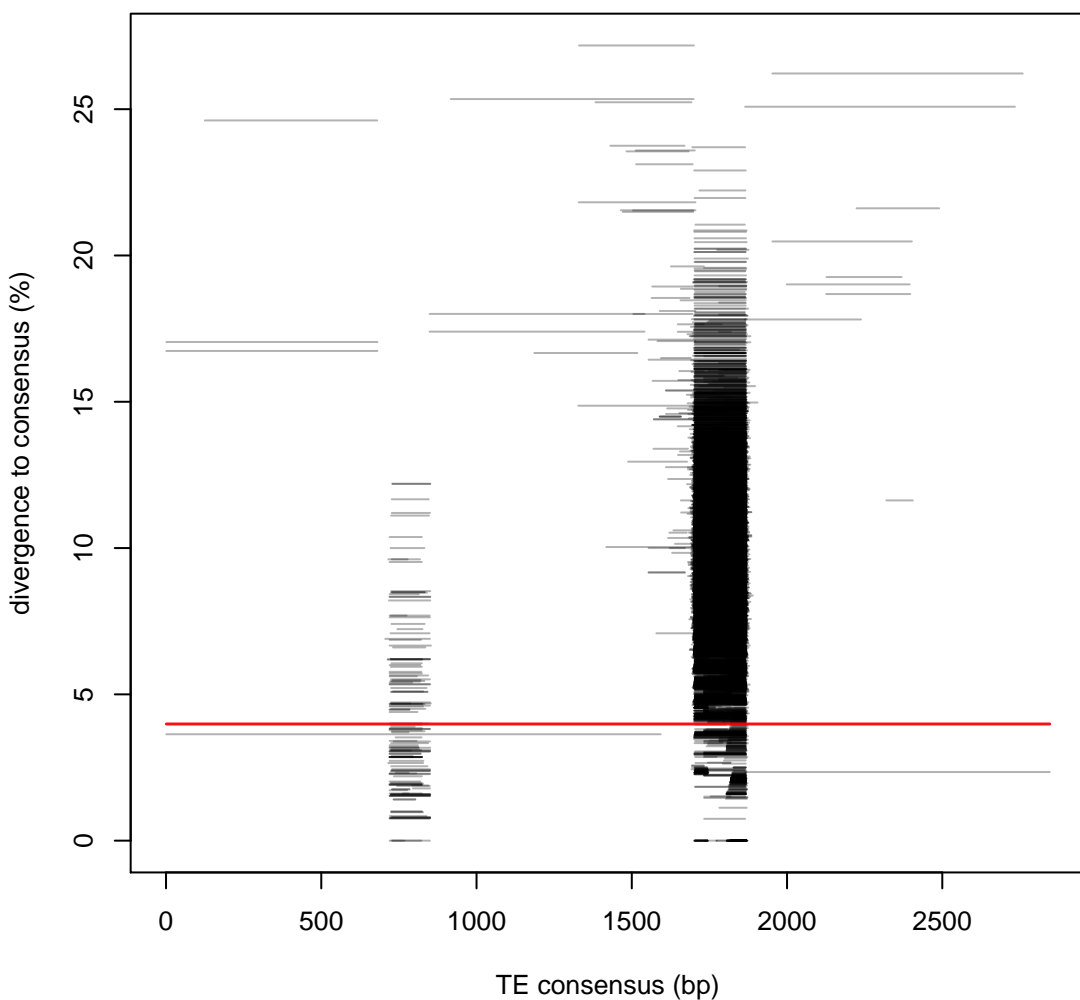
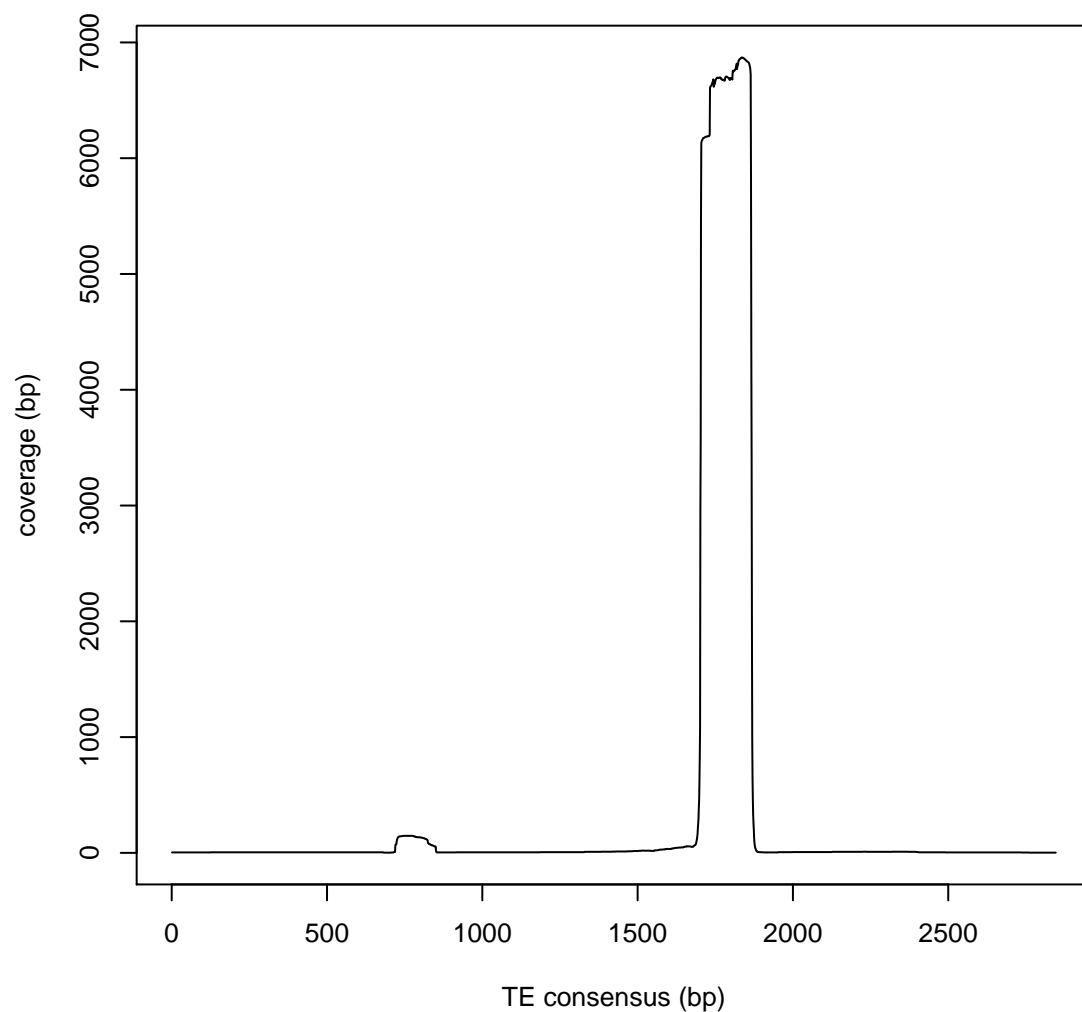


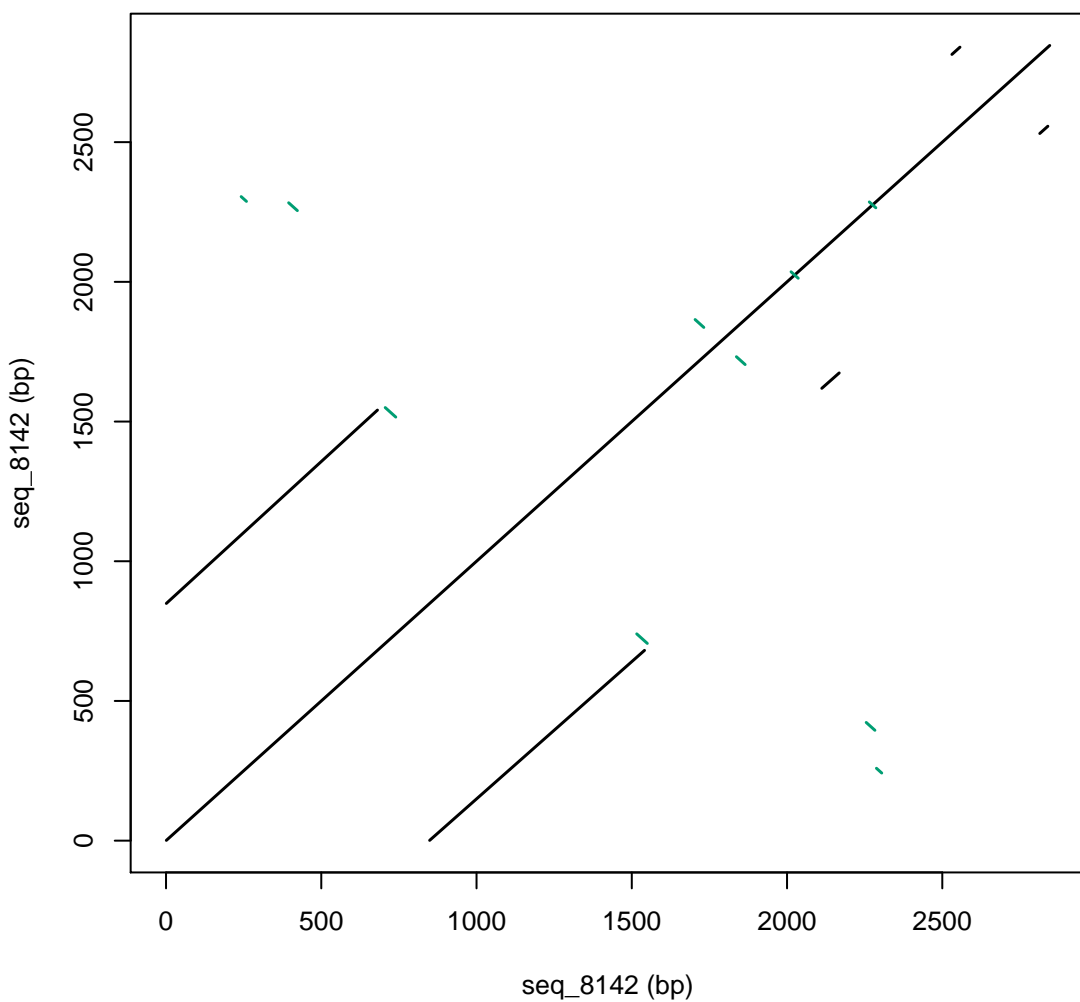
TE: seq_8142
consensus size: 2846bp; fragments: 7565; full length: 1 (≥ 2561.4 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

