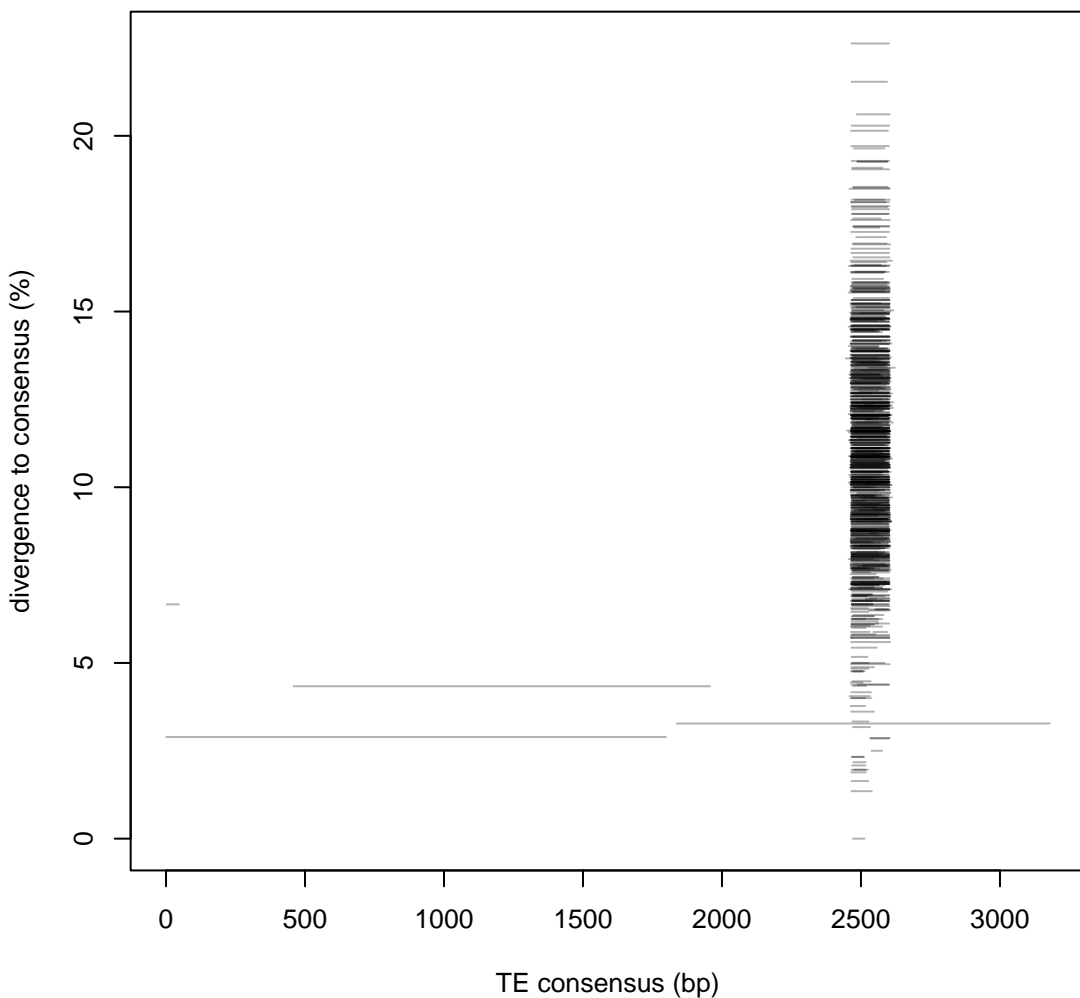
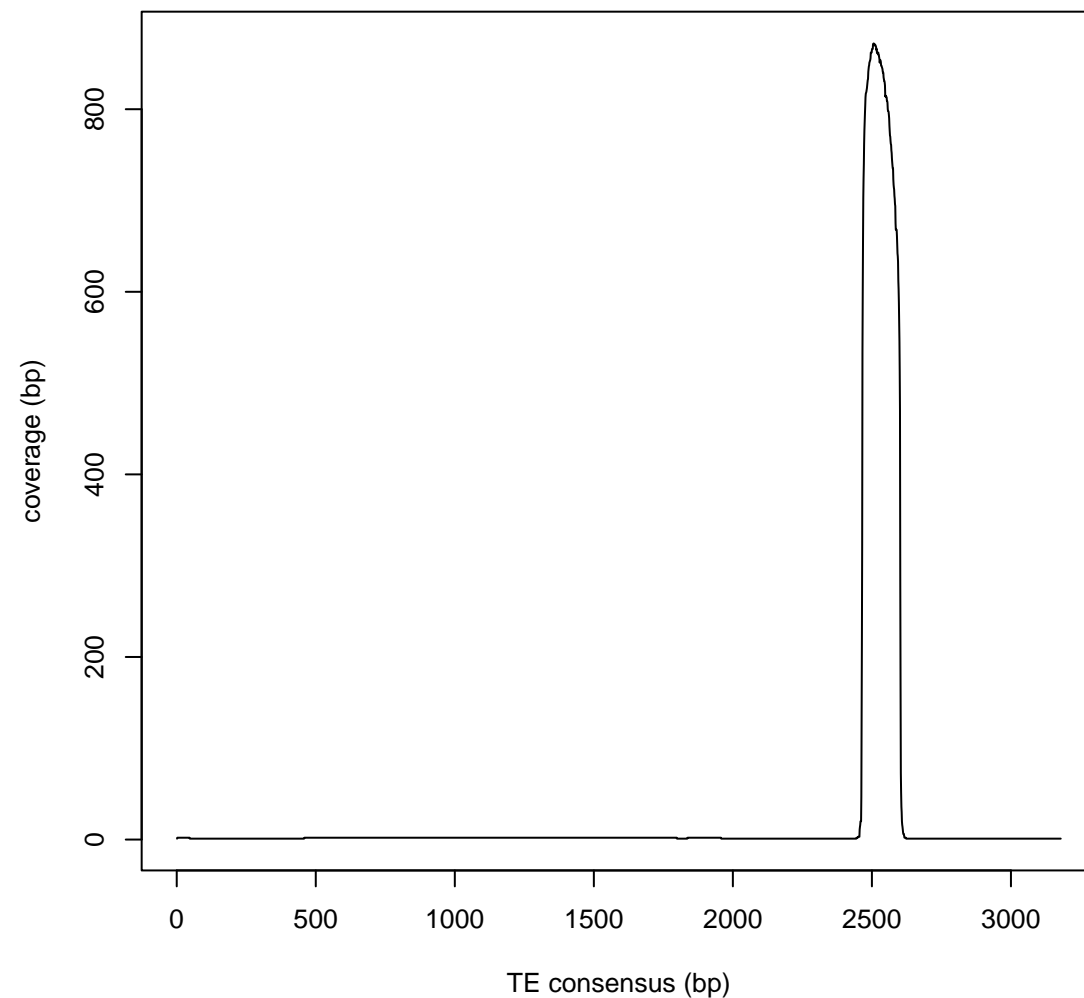


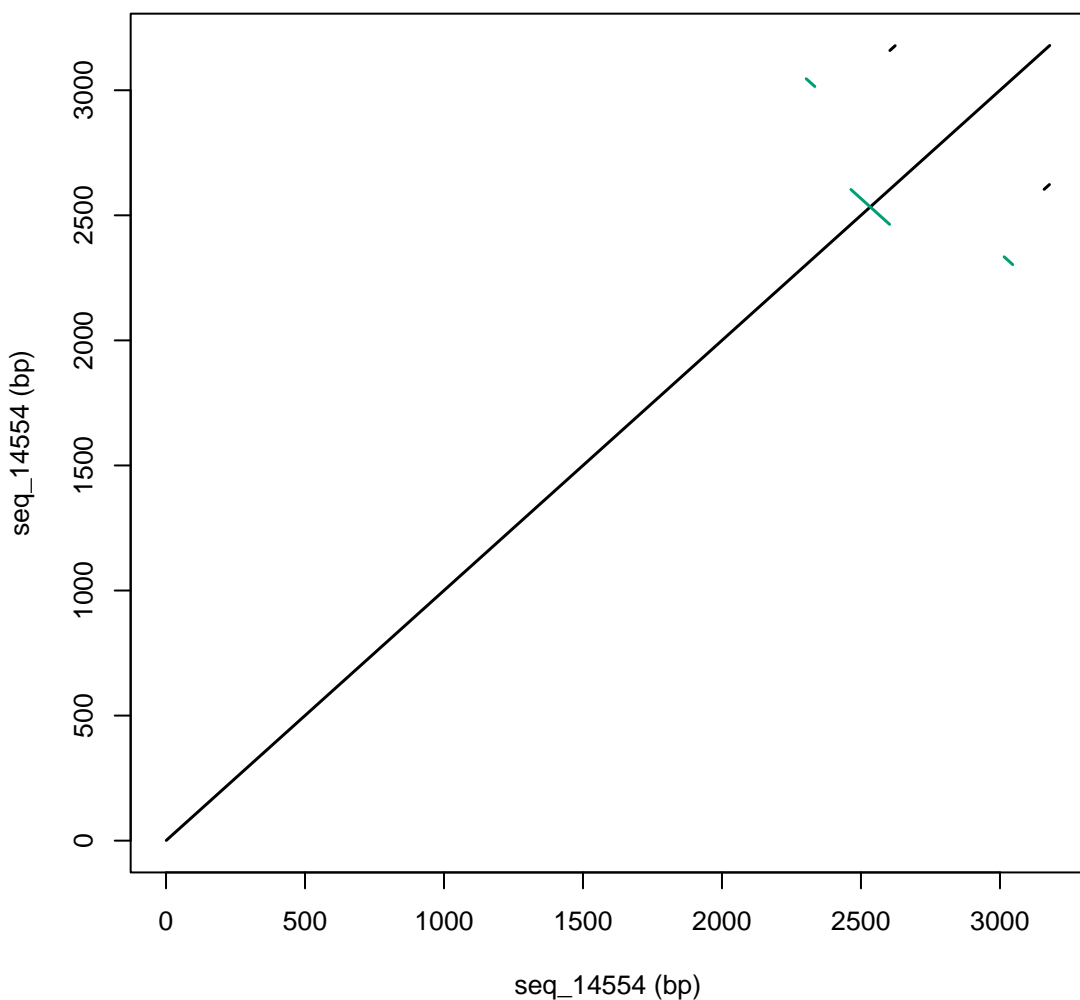
TE: seq_14554
consensus size: 3179bp; fragments: 900; full length: 0 (≥ 2861.1 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

