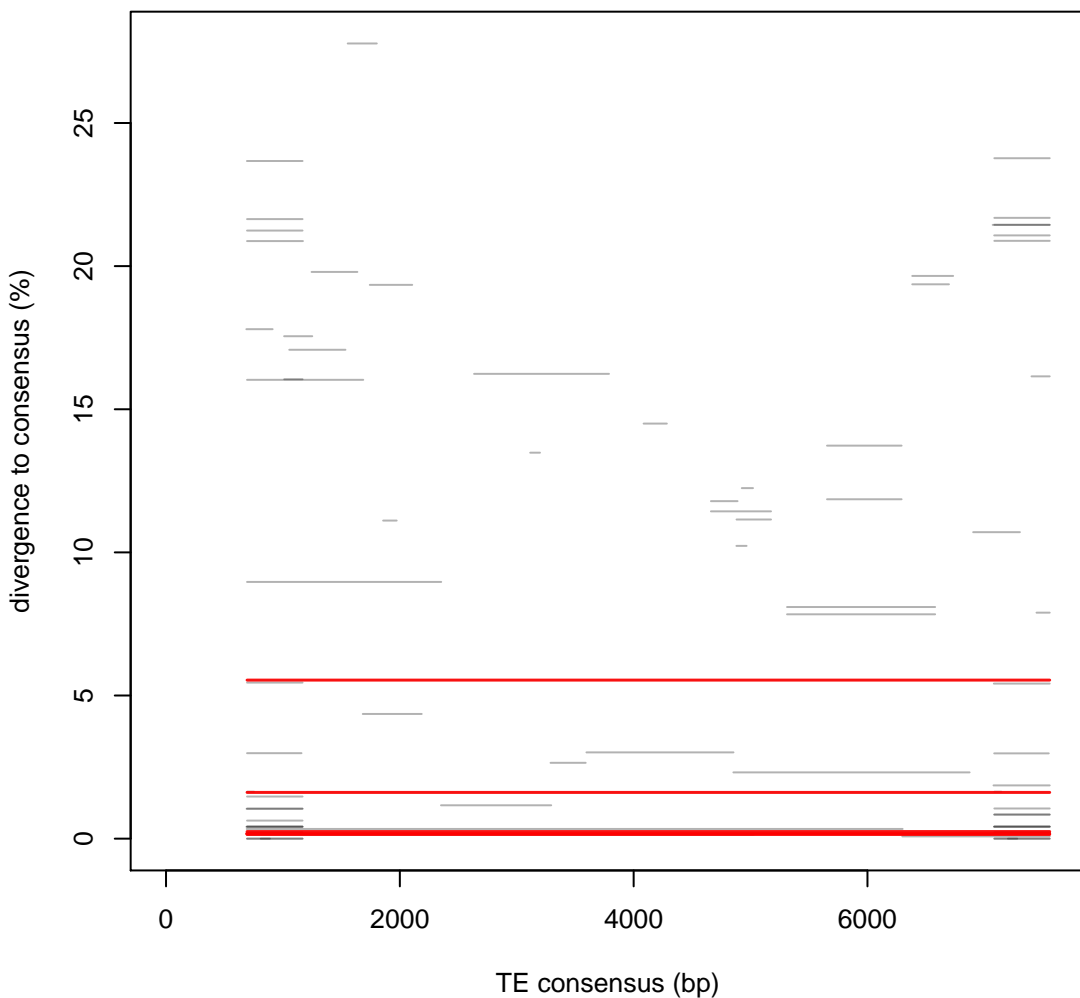
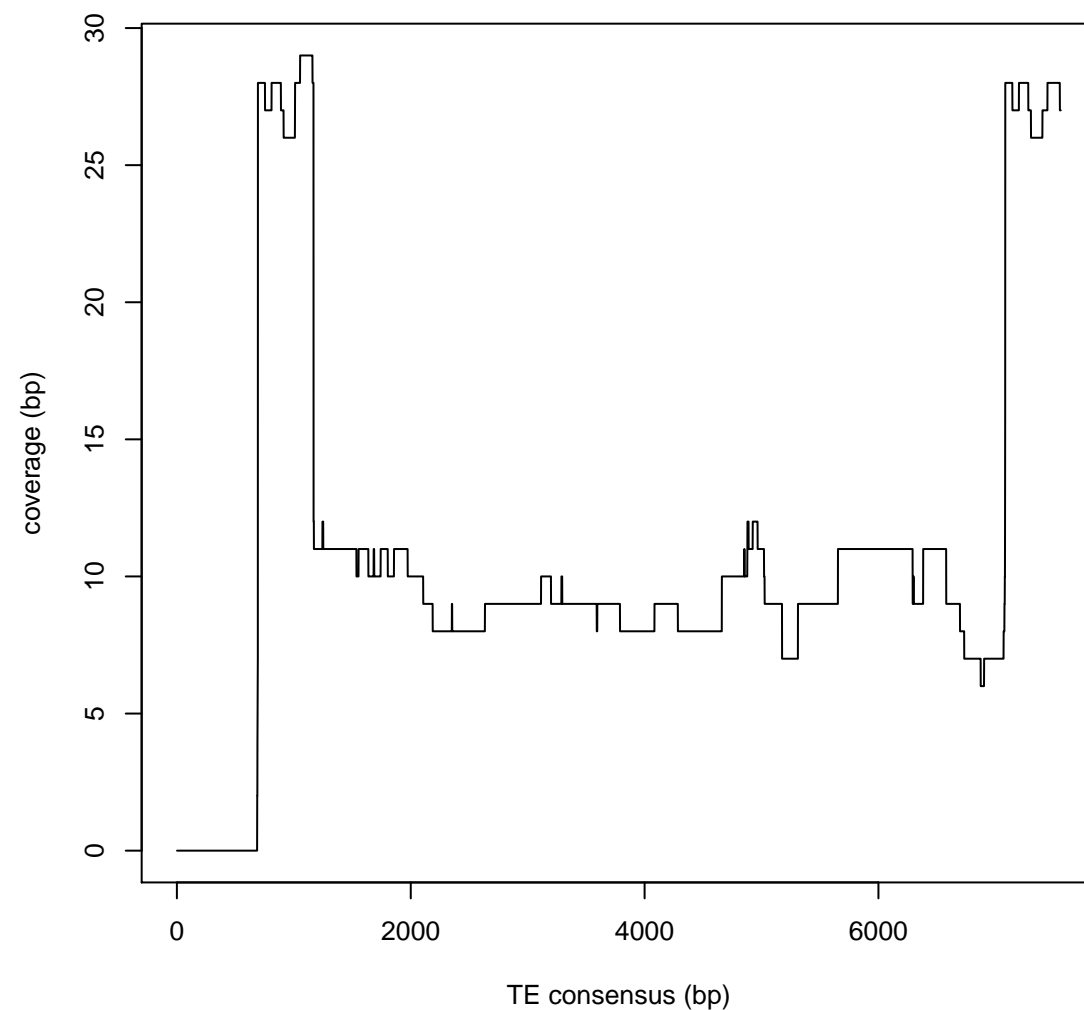


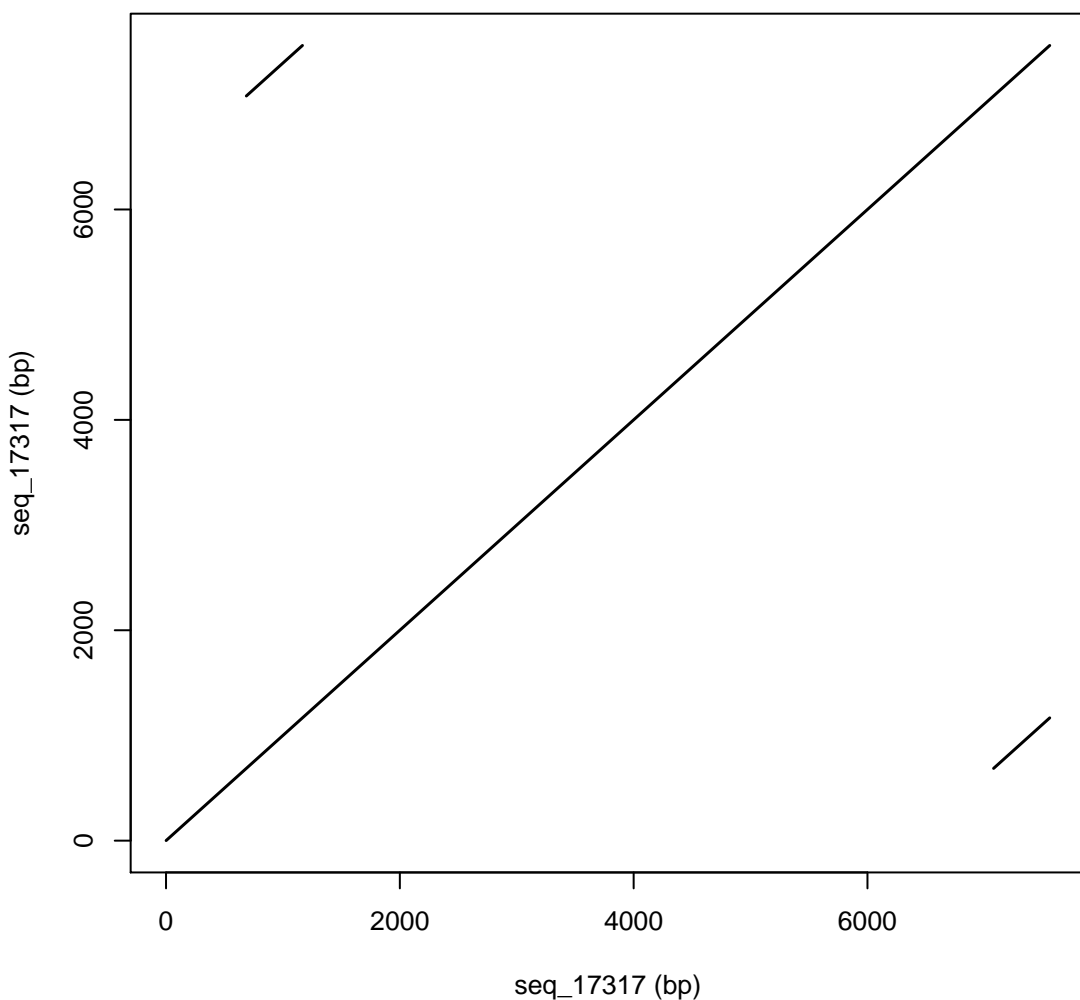
TE: seq_17317
consensus size: 7559bp; fragments: 81; full length: 5 (>=6803.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

