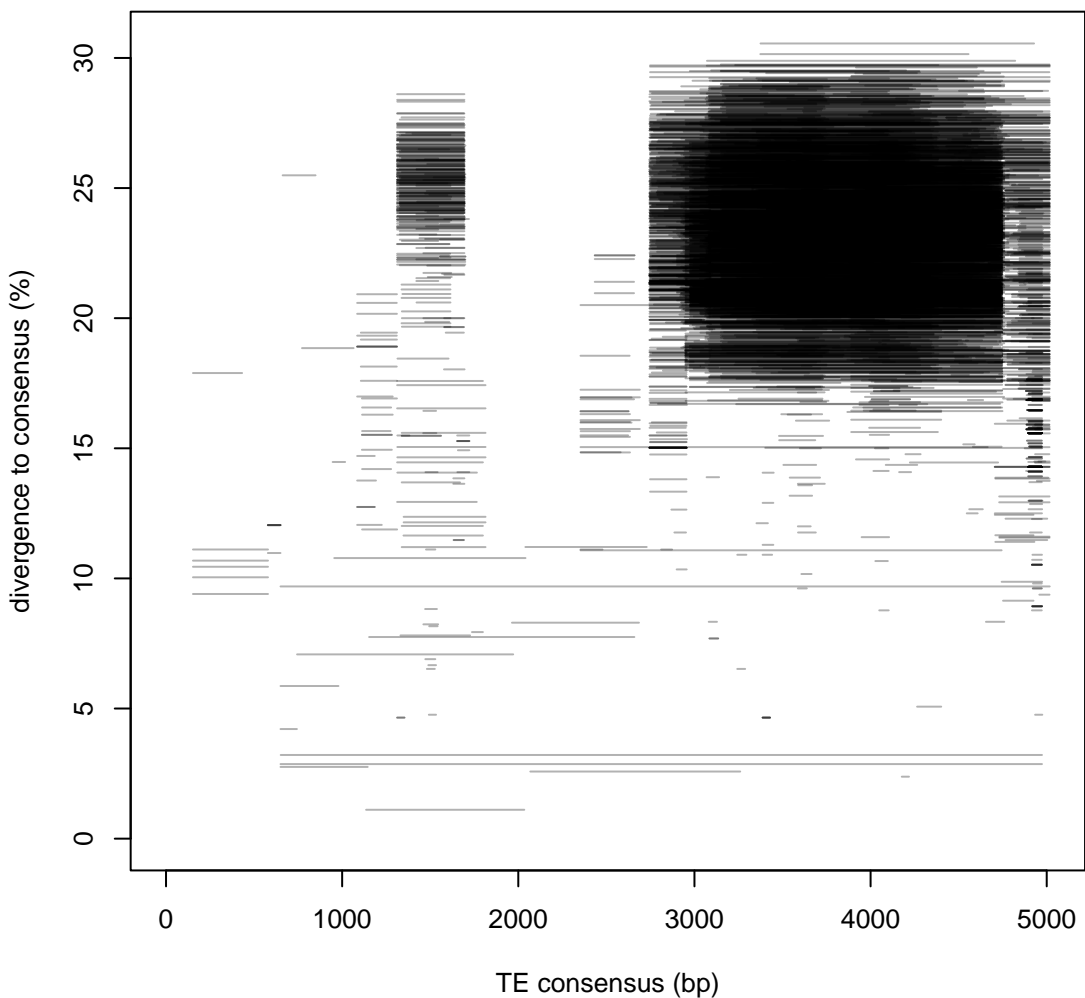
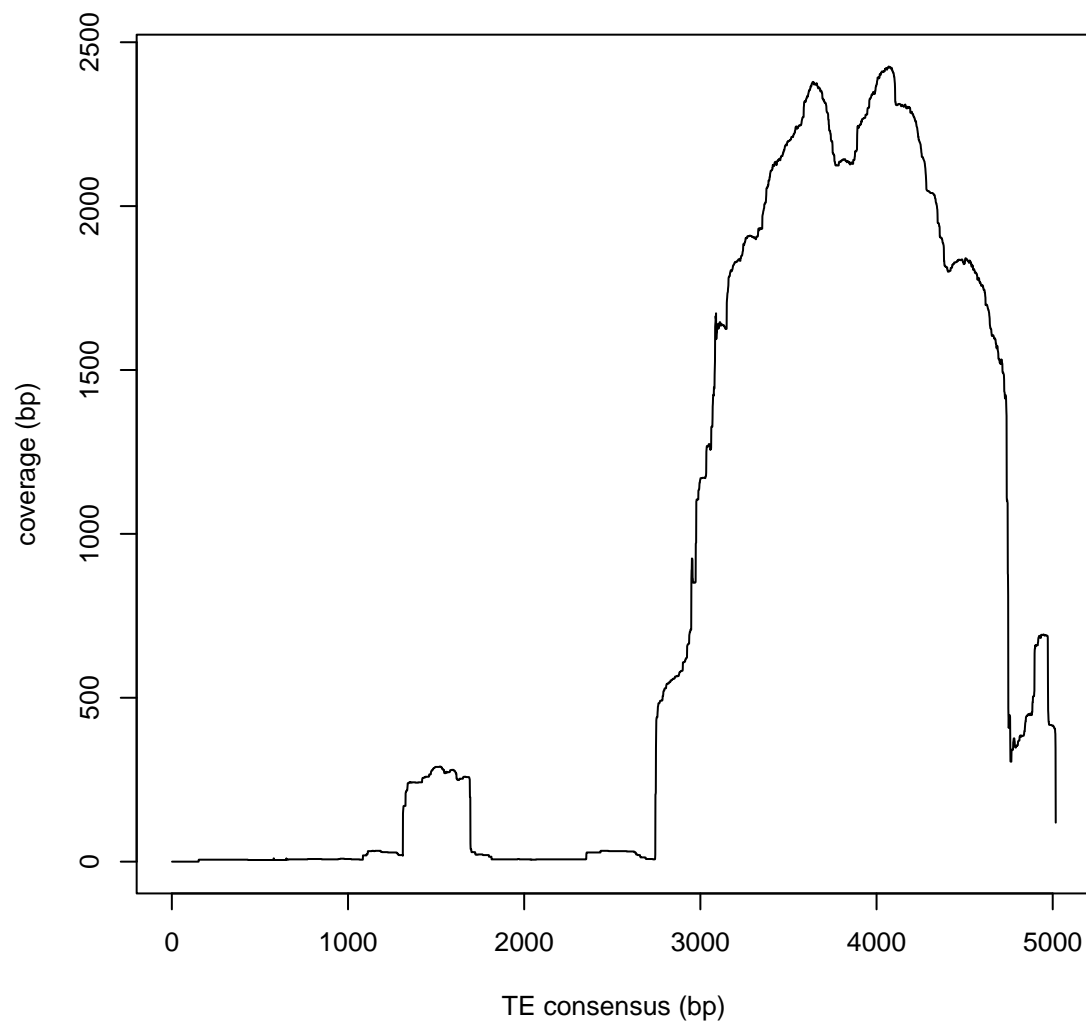


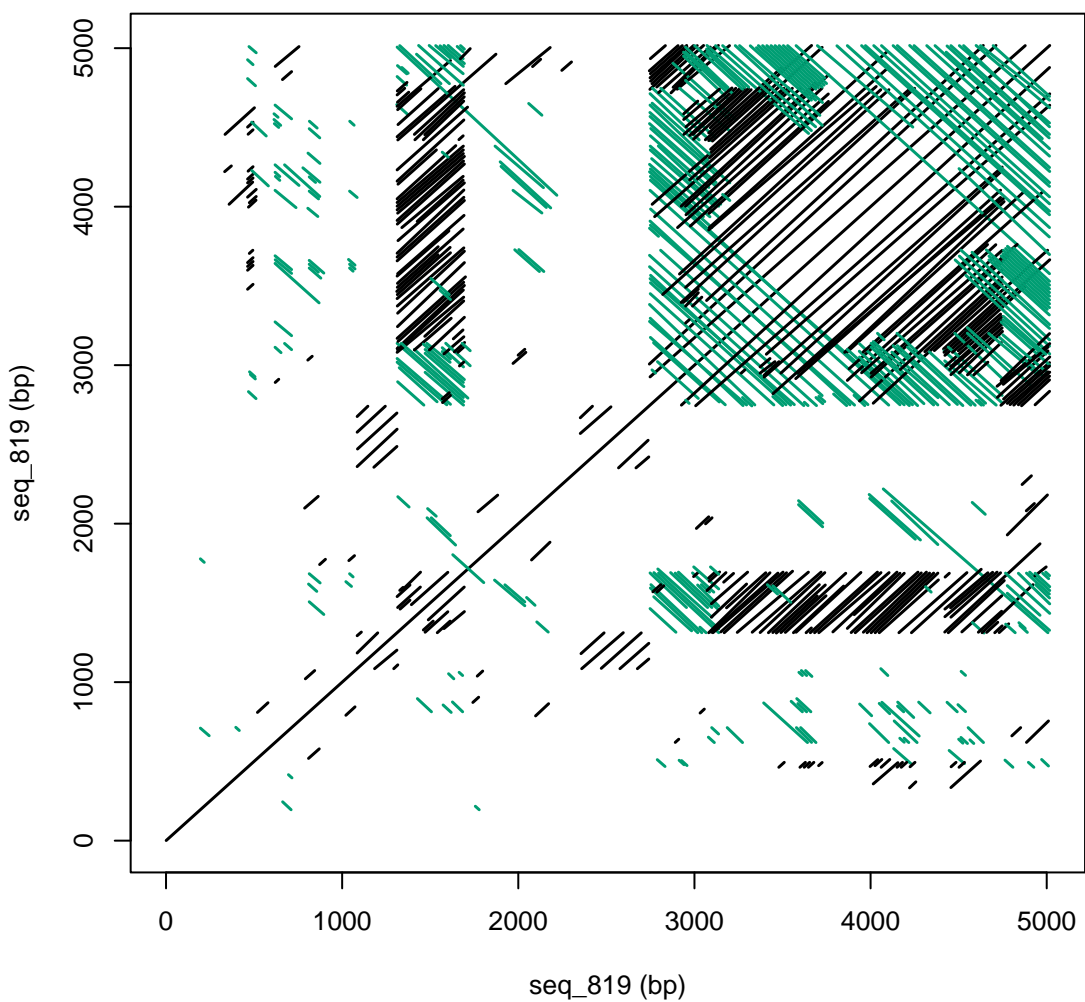
TE: seq_819
consensus size: 5017bp; fragments: 4212; full length: 0 (≥ 4515.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

