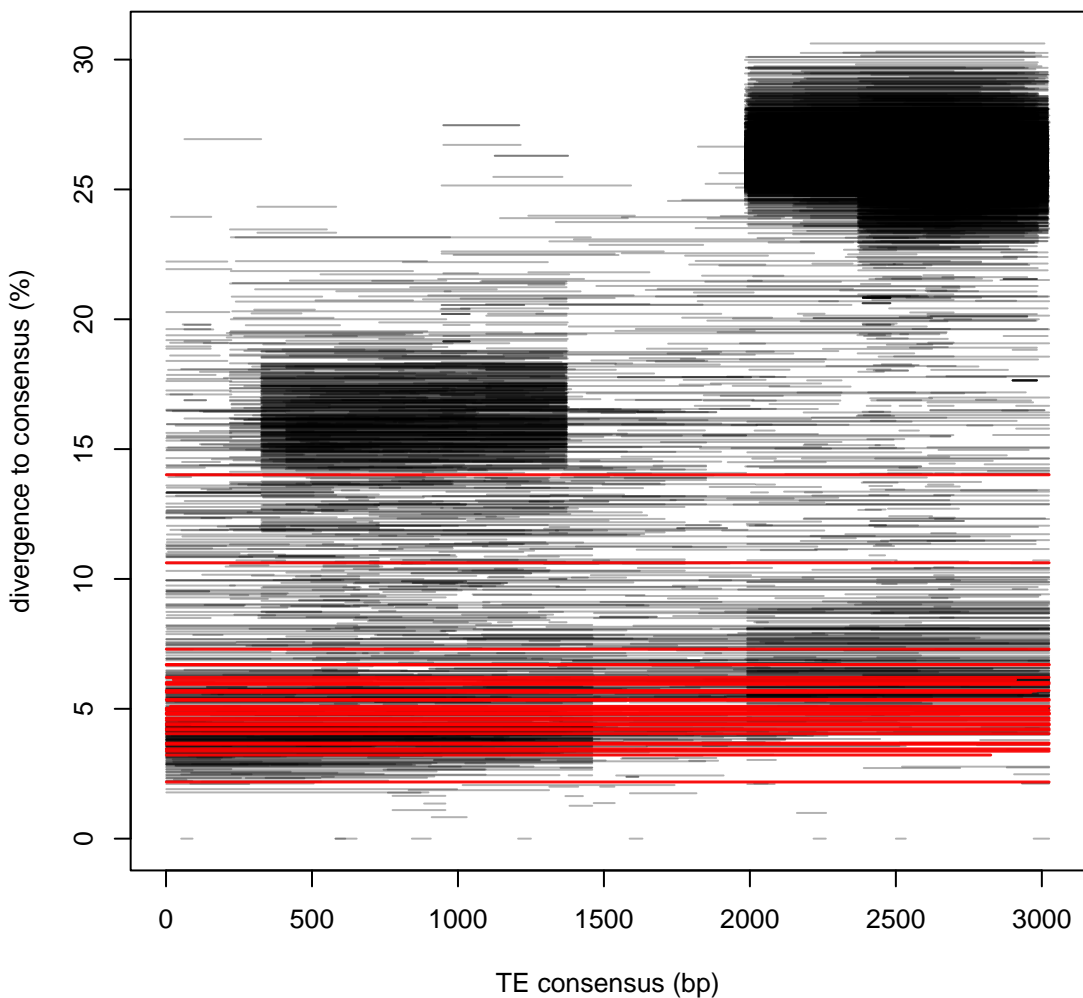
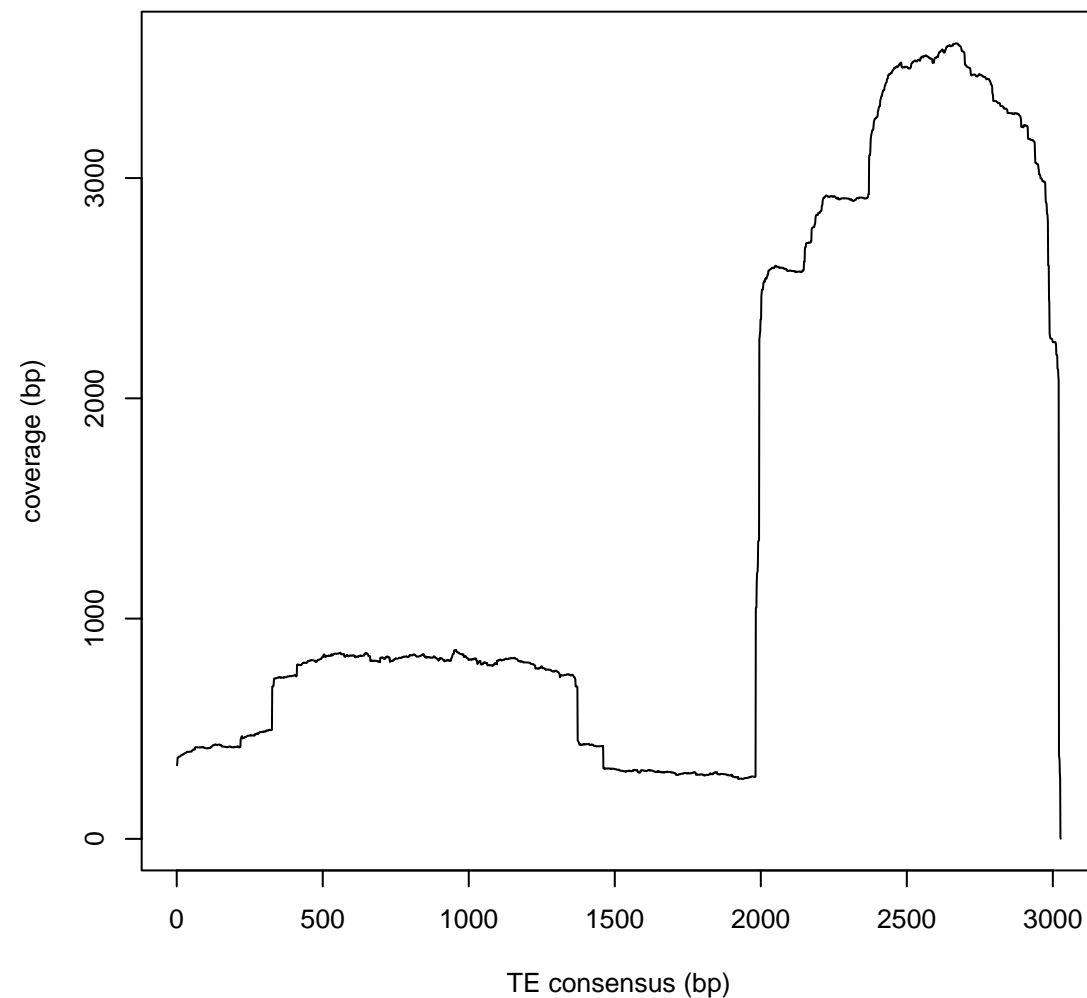


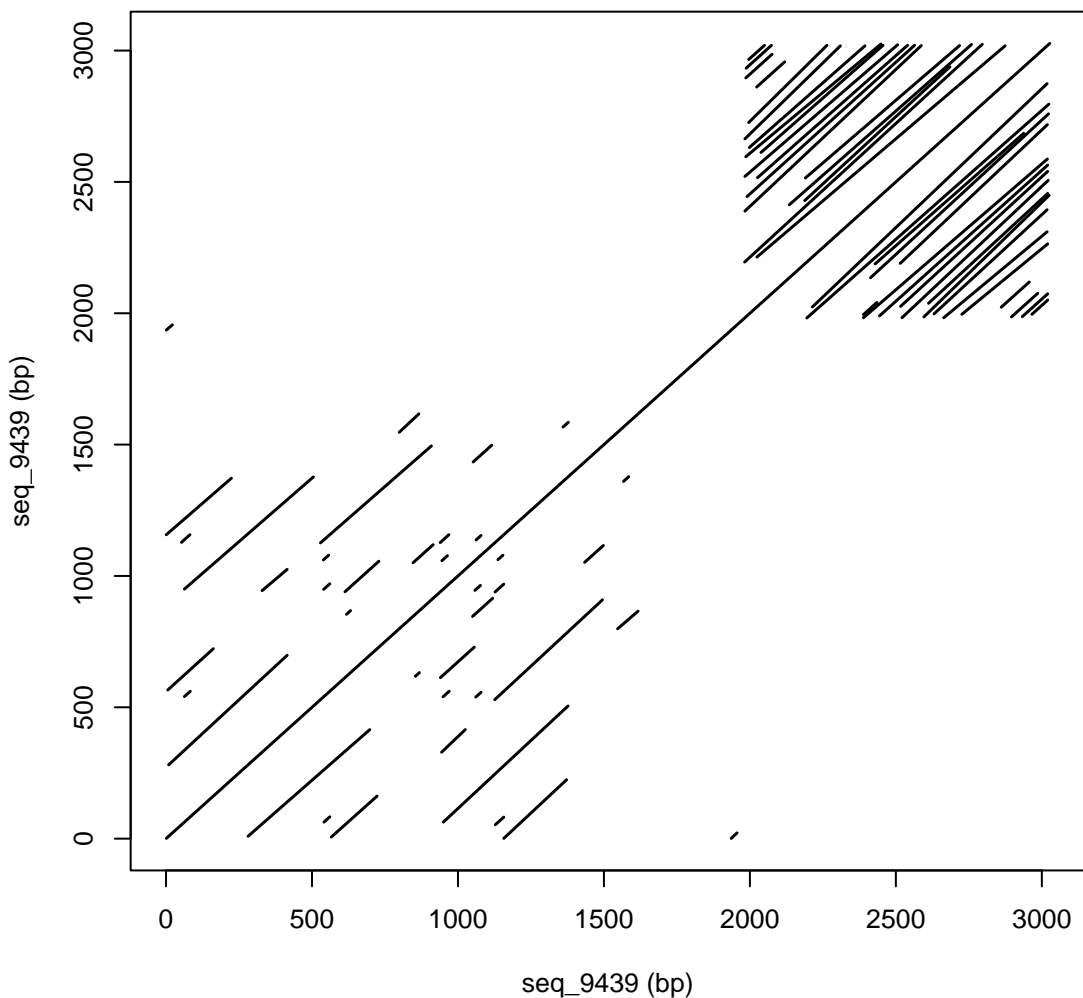
TE: seq_9439
consensus size: 3027bp; fragments: 5498; full length: 37 (>=2724.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

