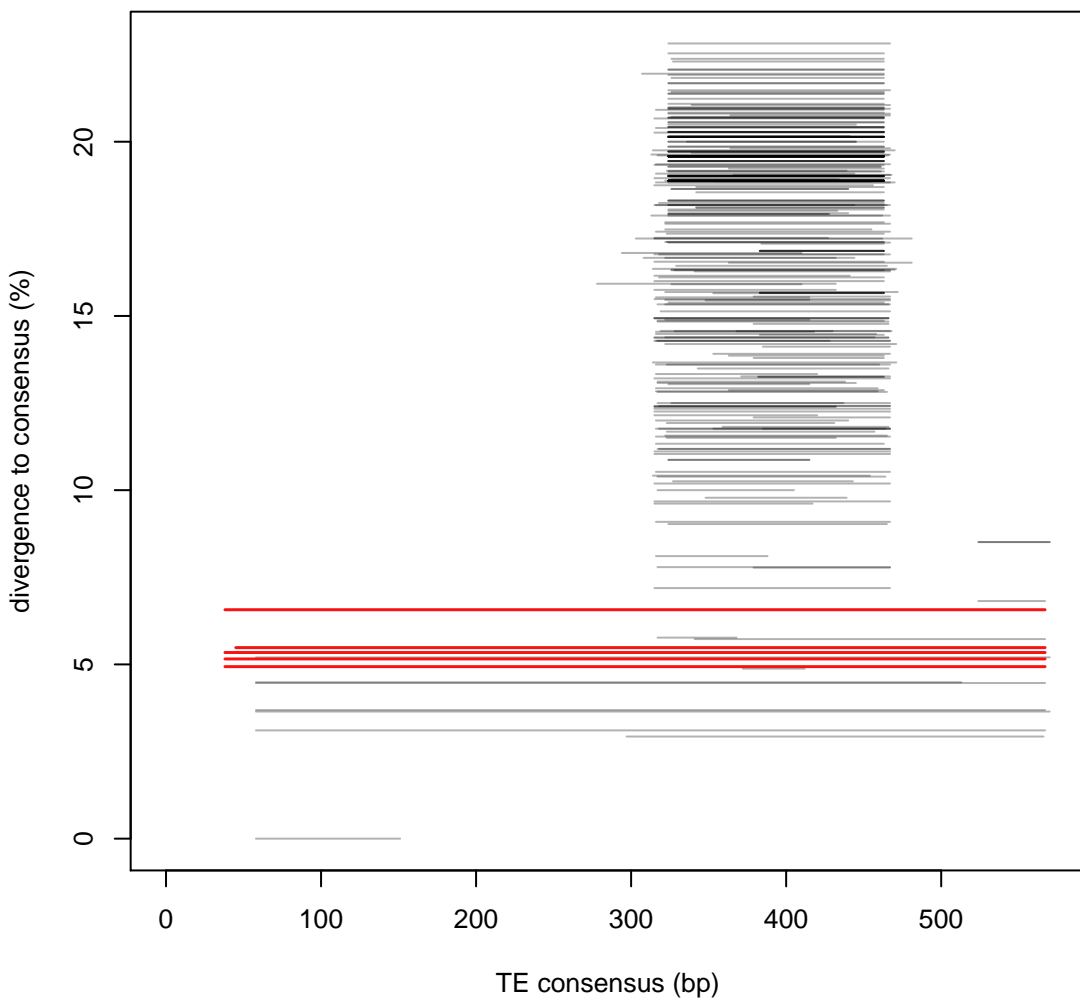
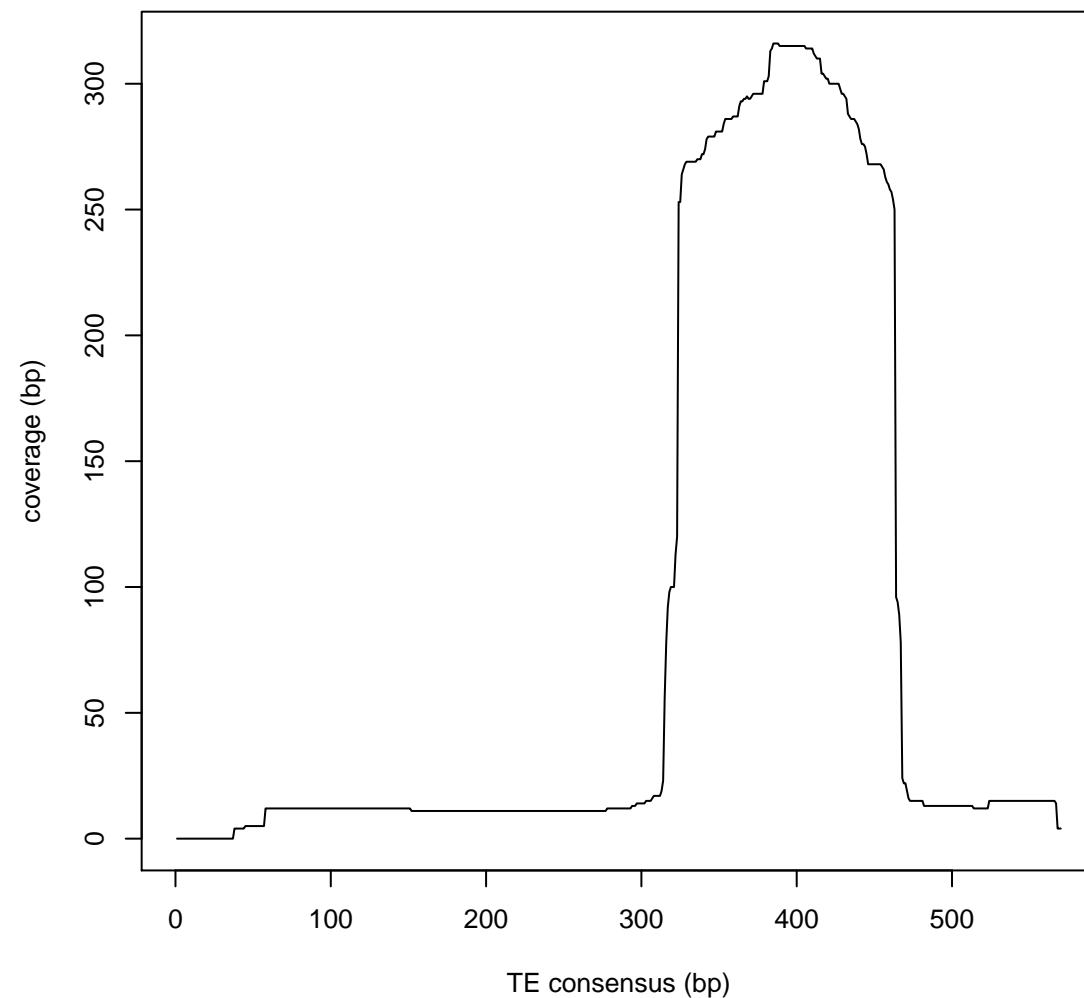


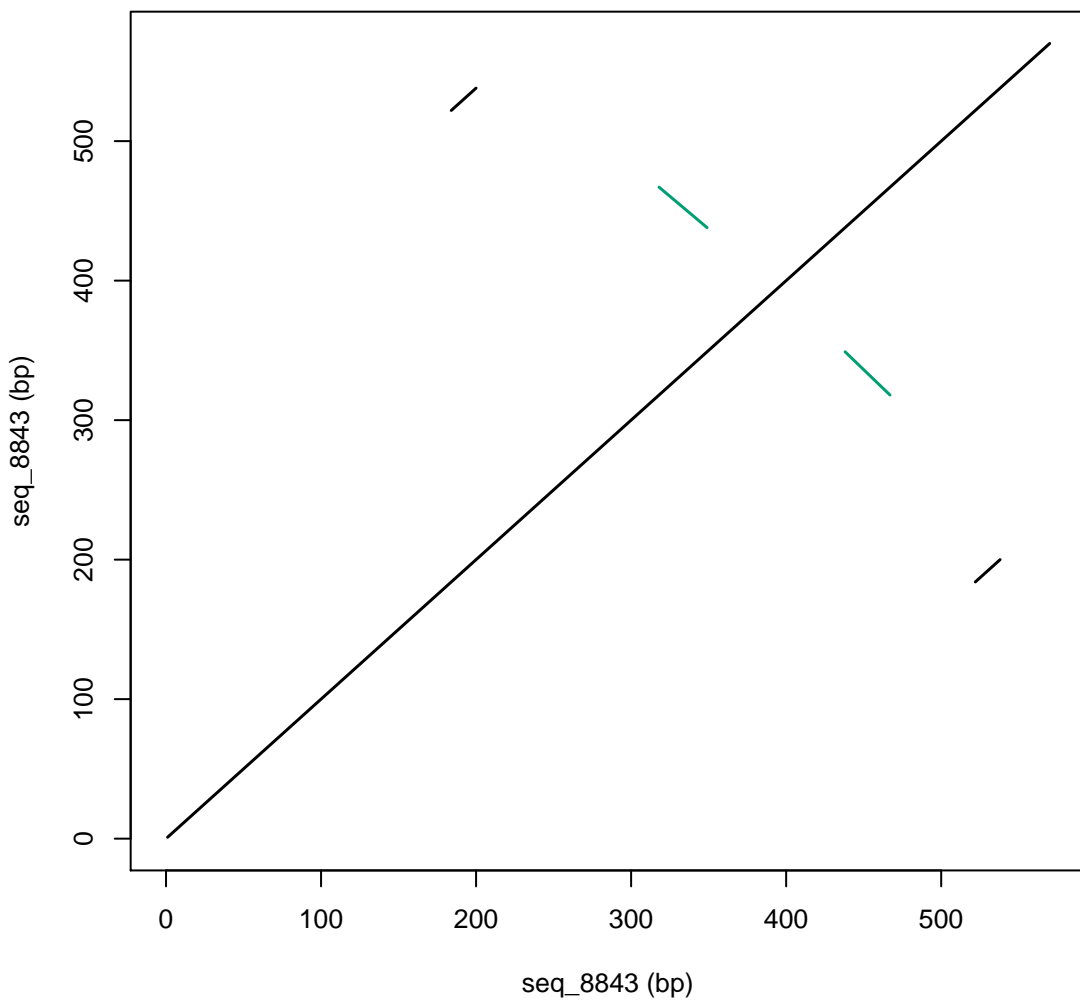
TE: seq_8843
consensus size: 570bp; fragments: 321; full length: 5 (>=513bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

