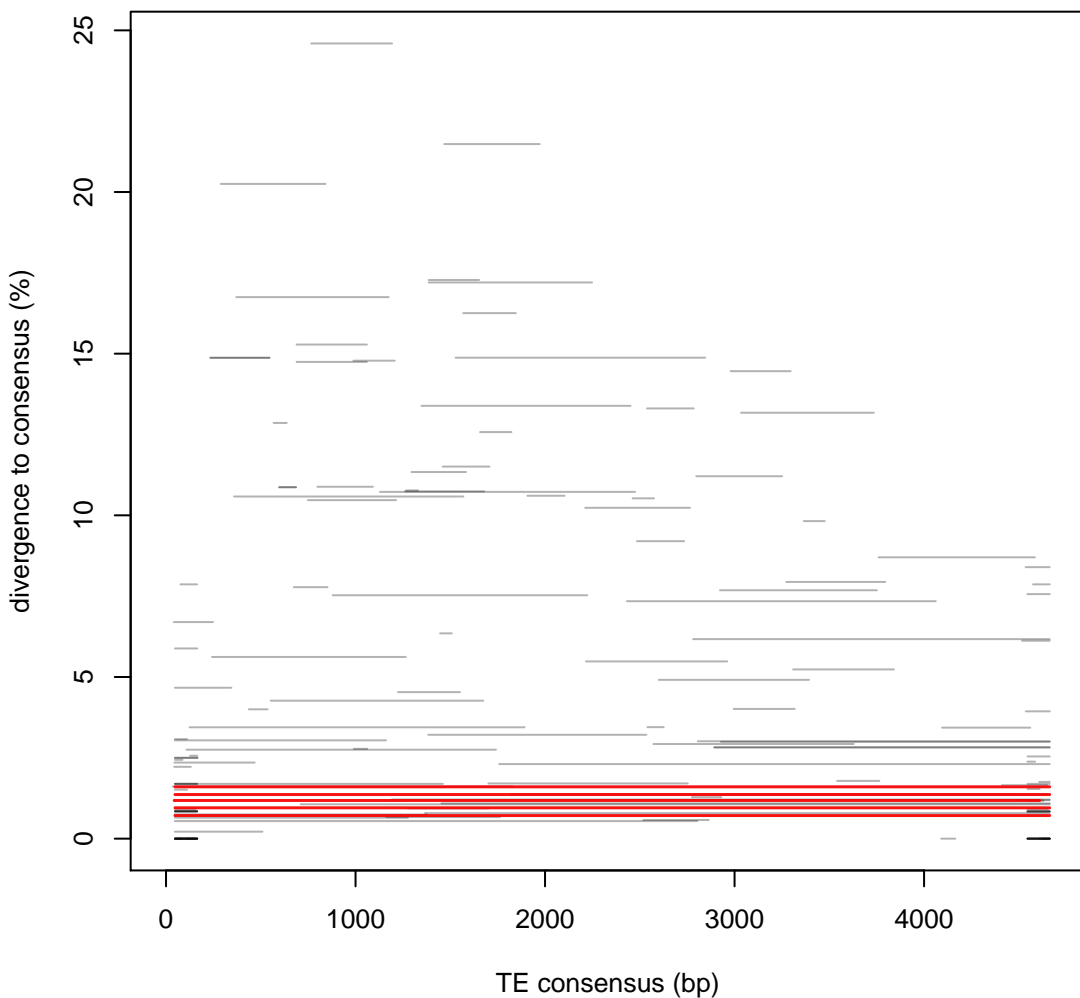
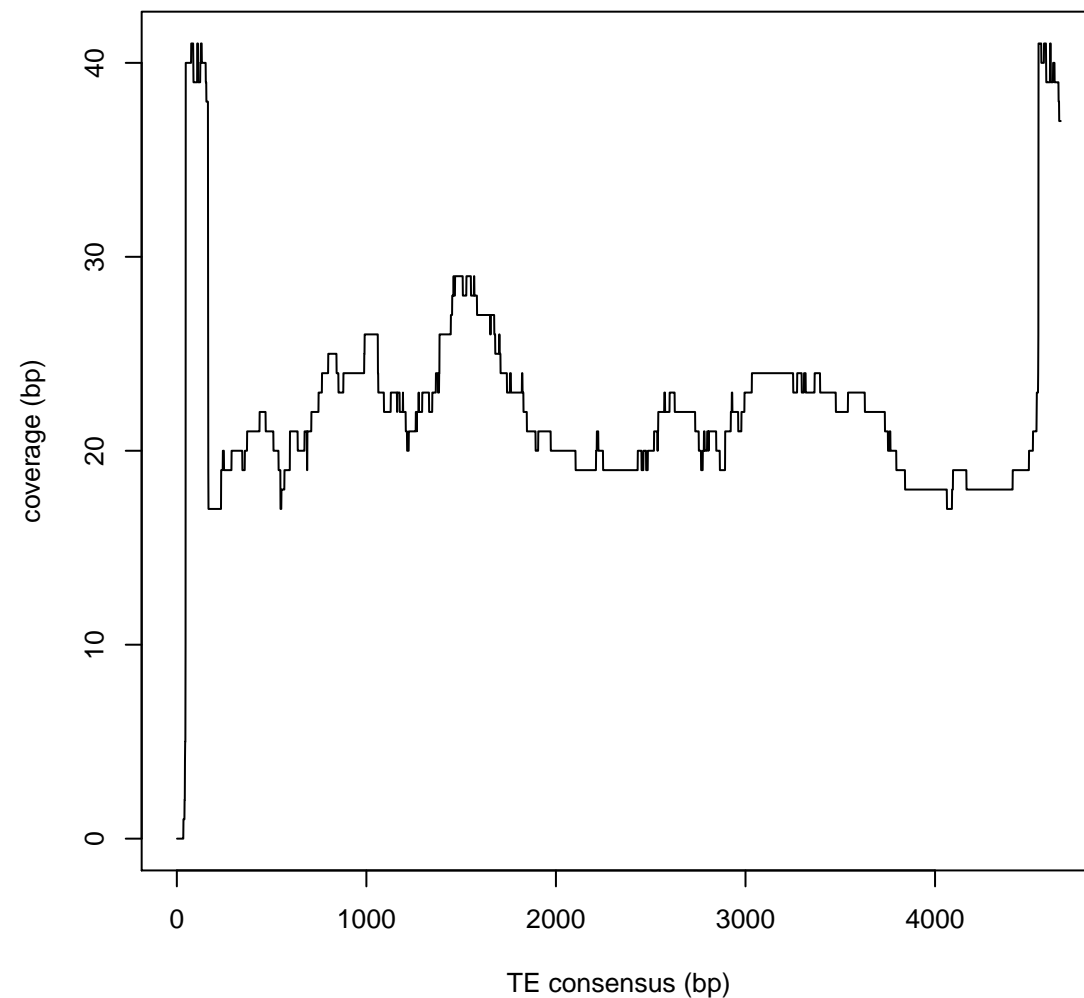


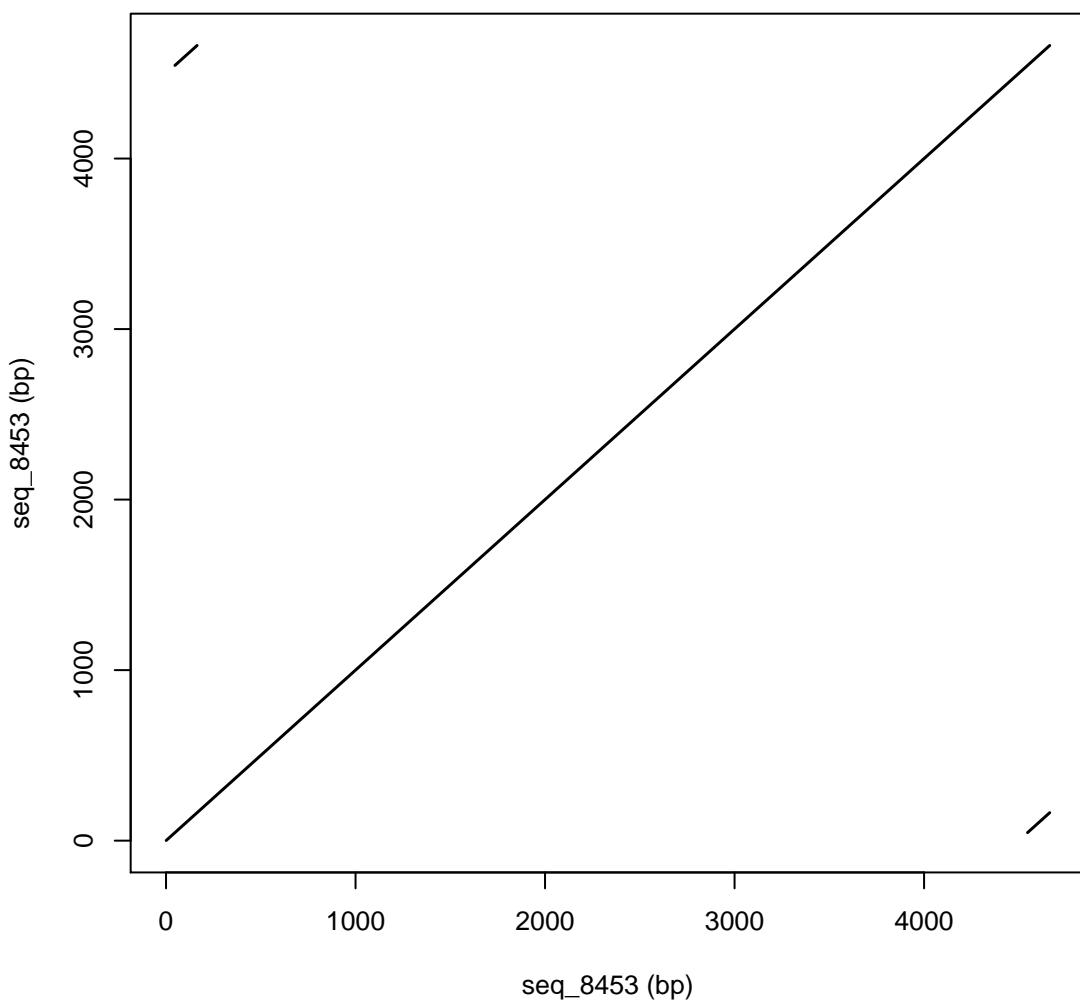
TE: seq_8453
consensus size: 4663bp; fragments: 145; full length: 5 (≥ 4196.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

