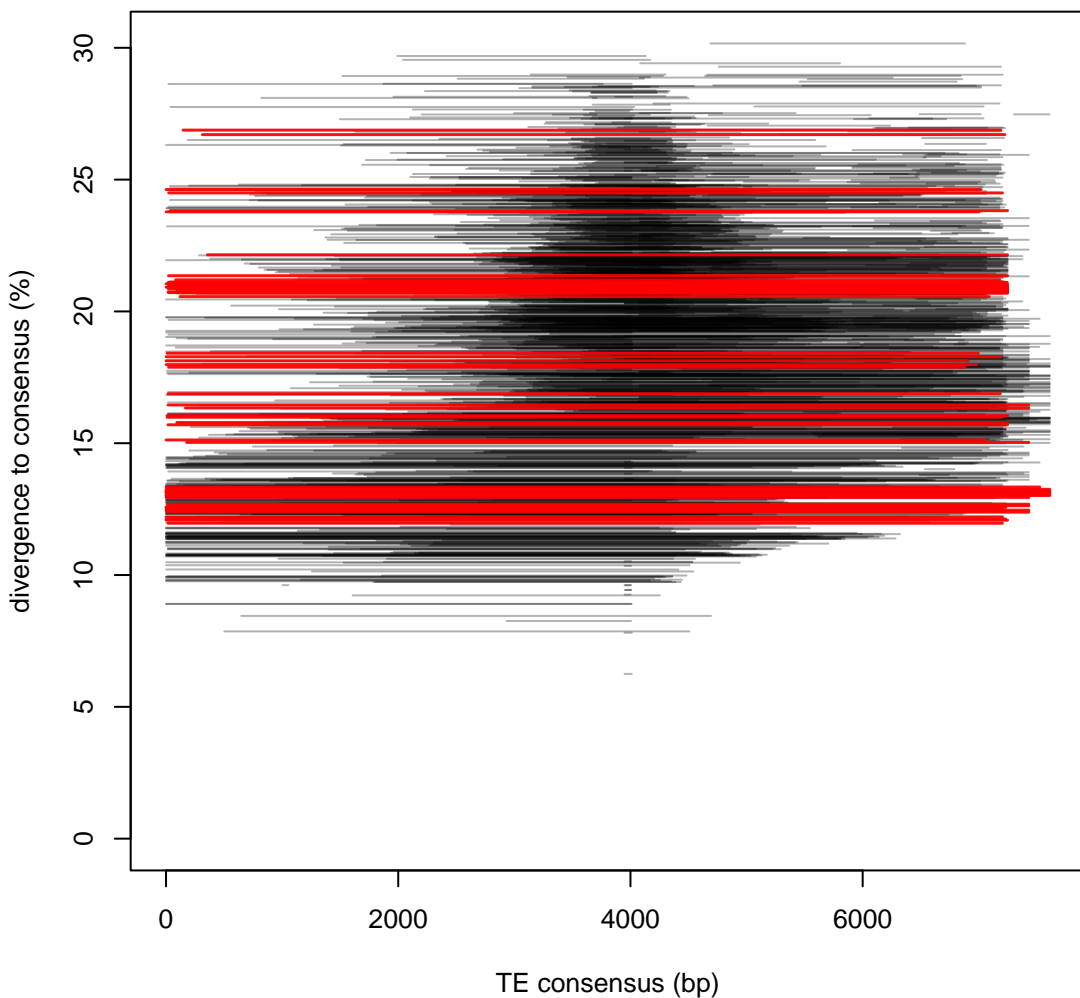
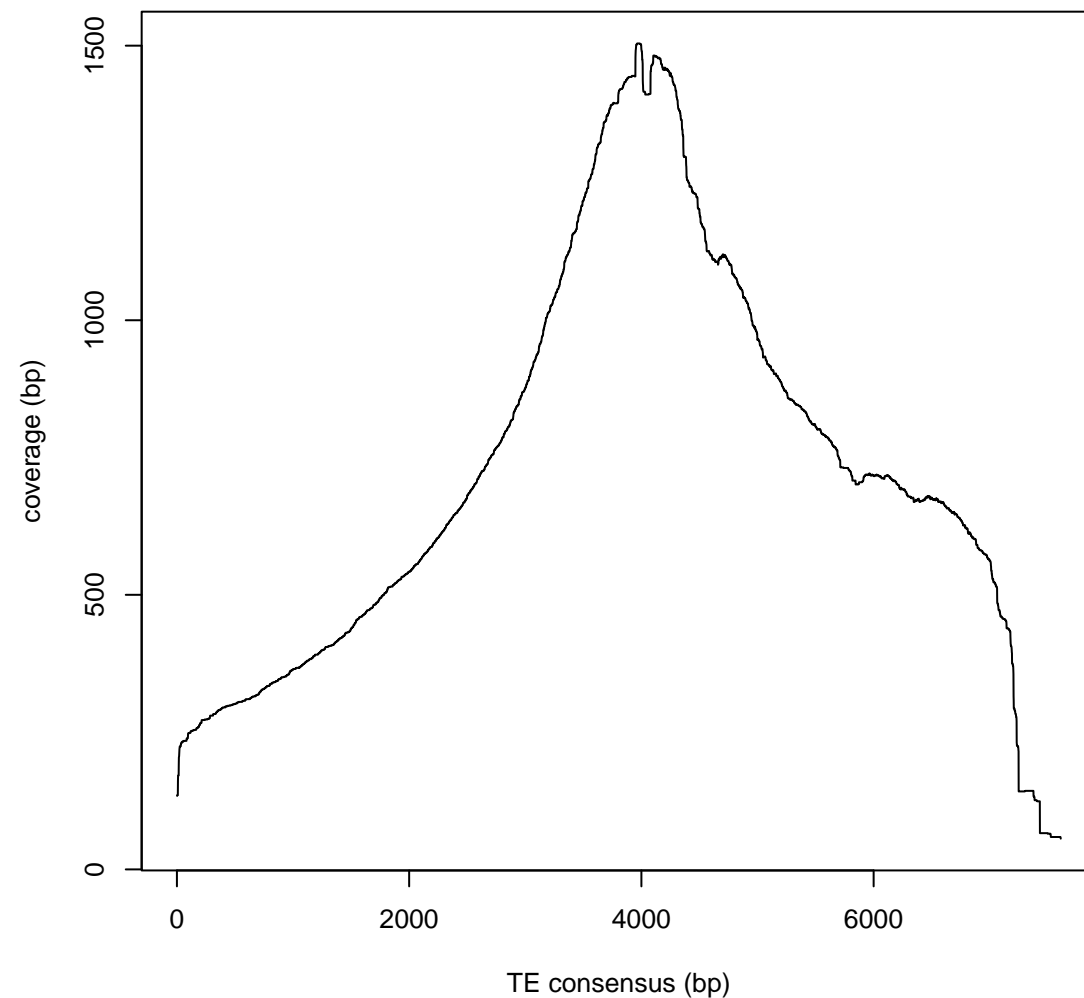


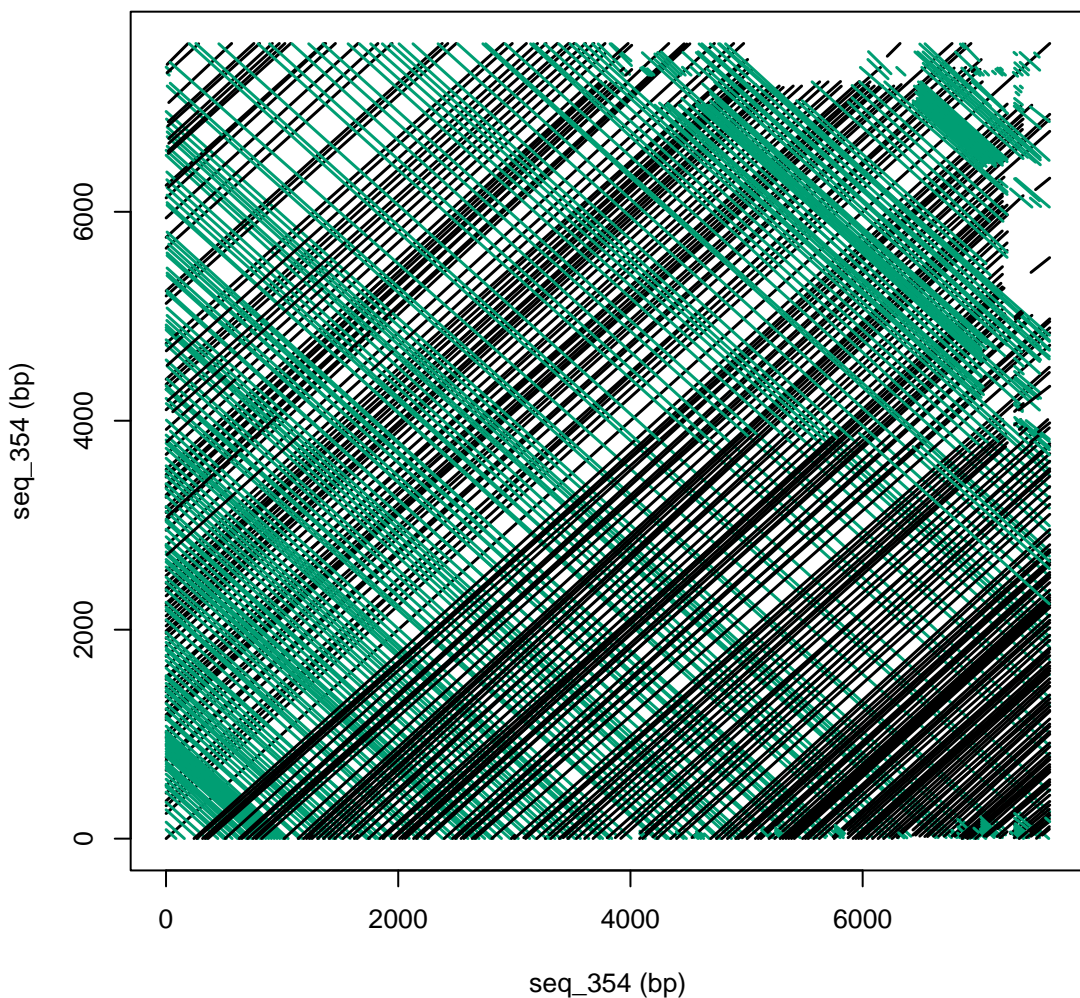
TE: seq_354
consensus size: 7611bp; fragments: 1803; full length: 80 (≥ 6849.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

