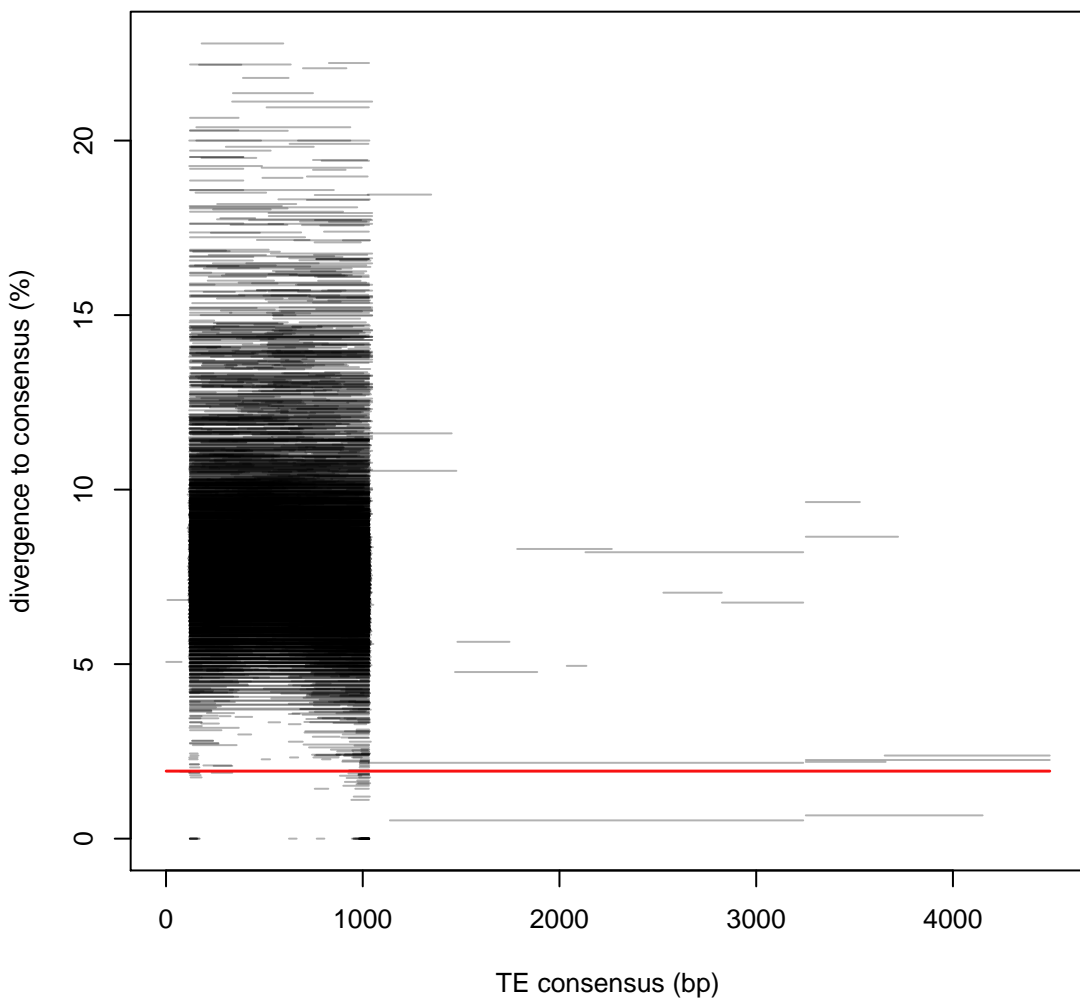
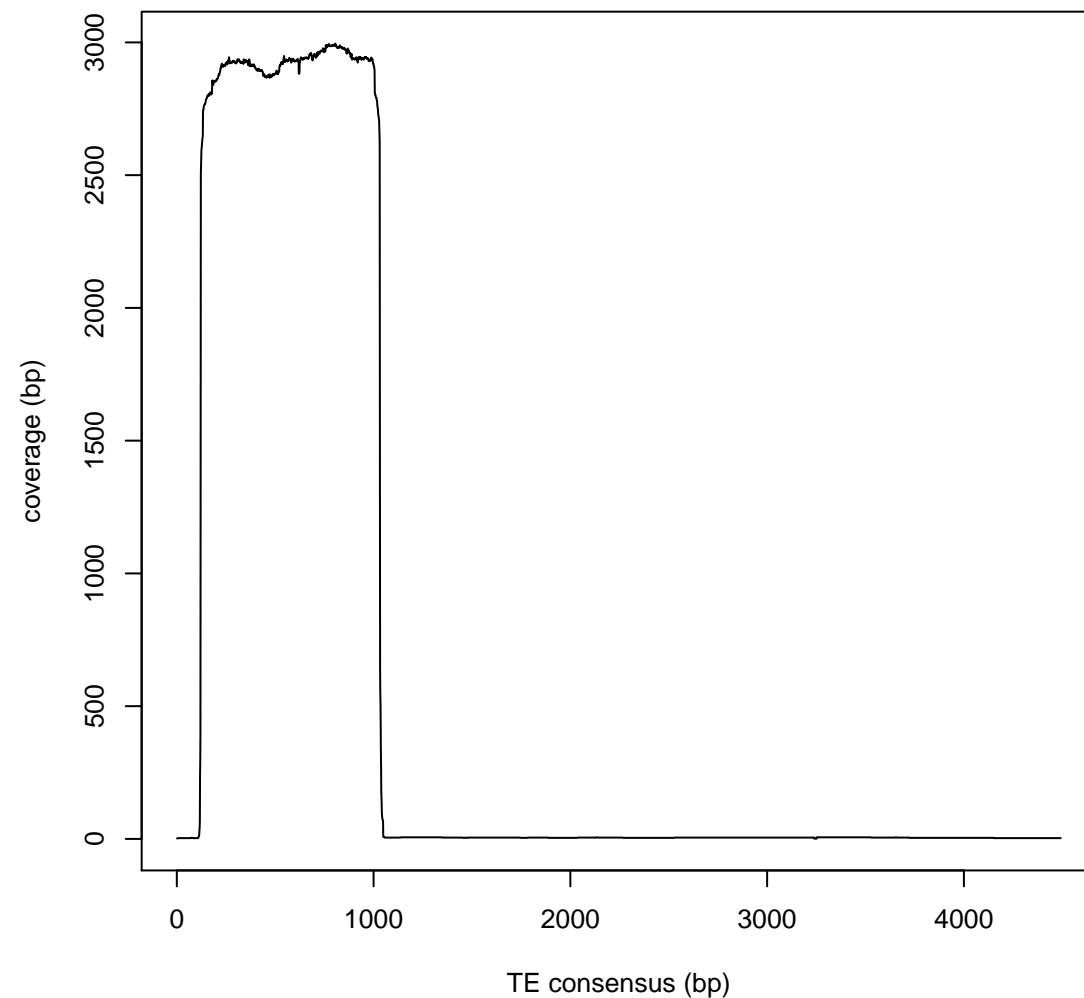


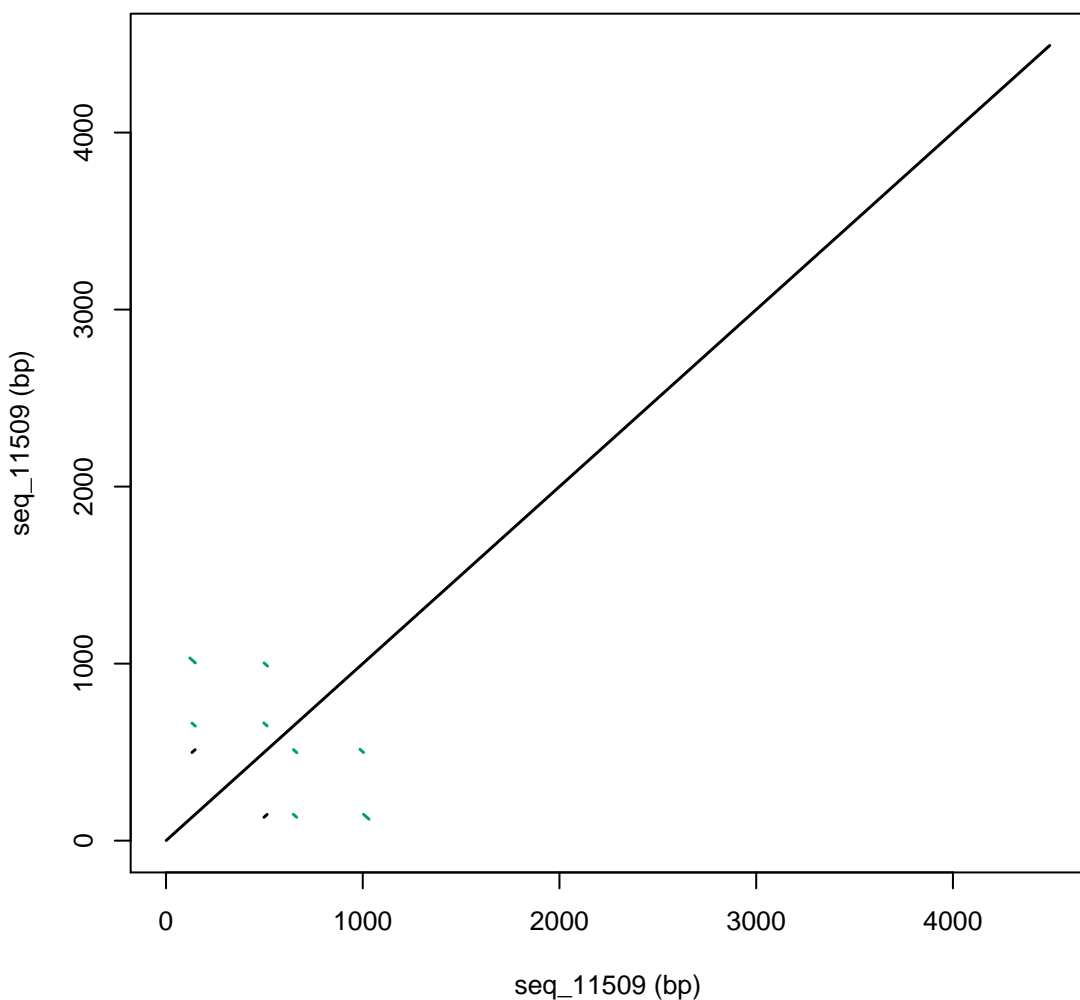
TE: seq_11509
consensus size: 4492bp; fragments: 4692; full length: 1 (≥ 4042.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

