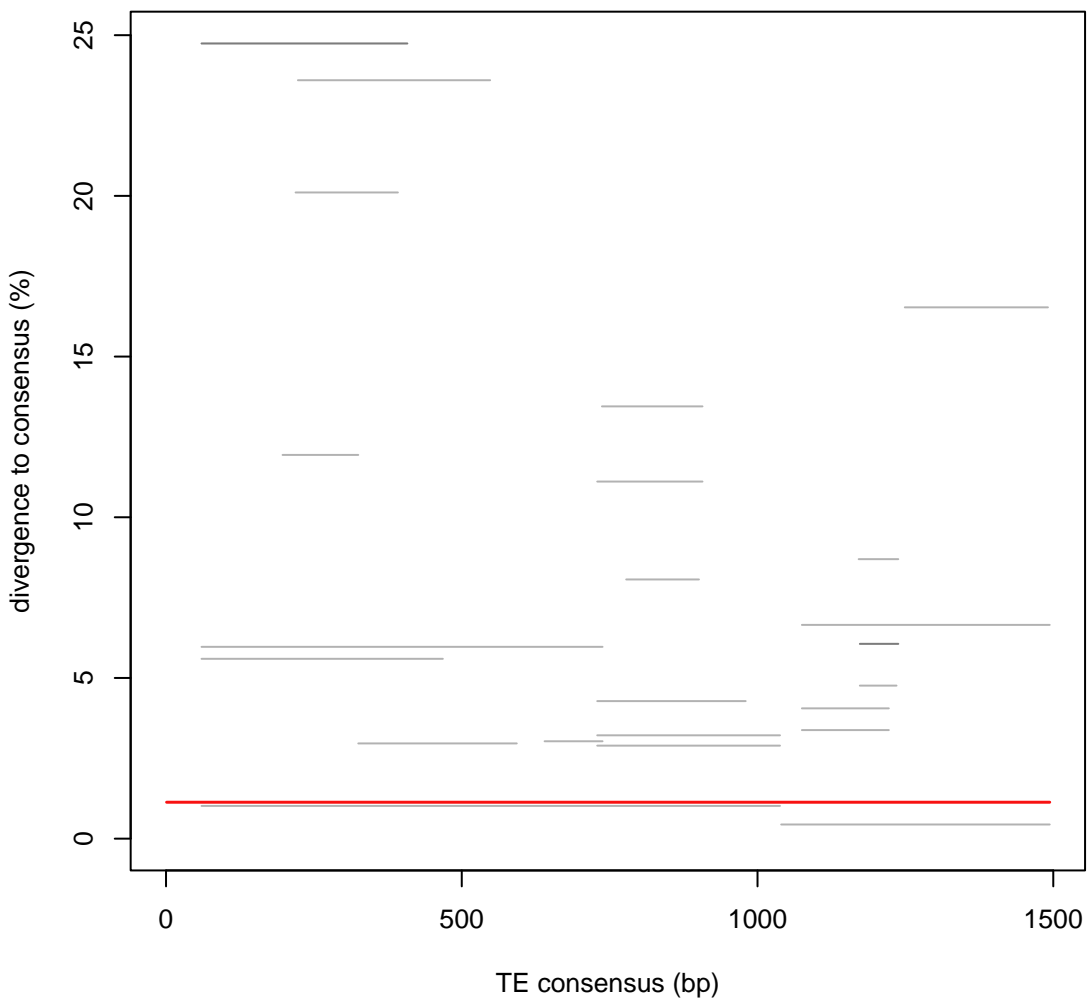
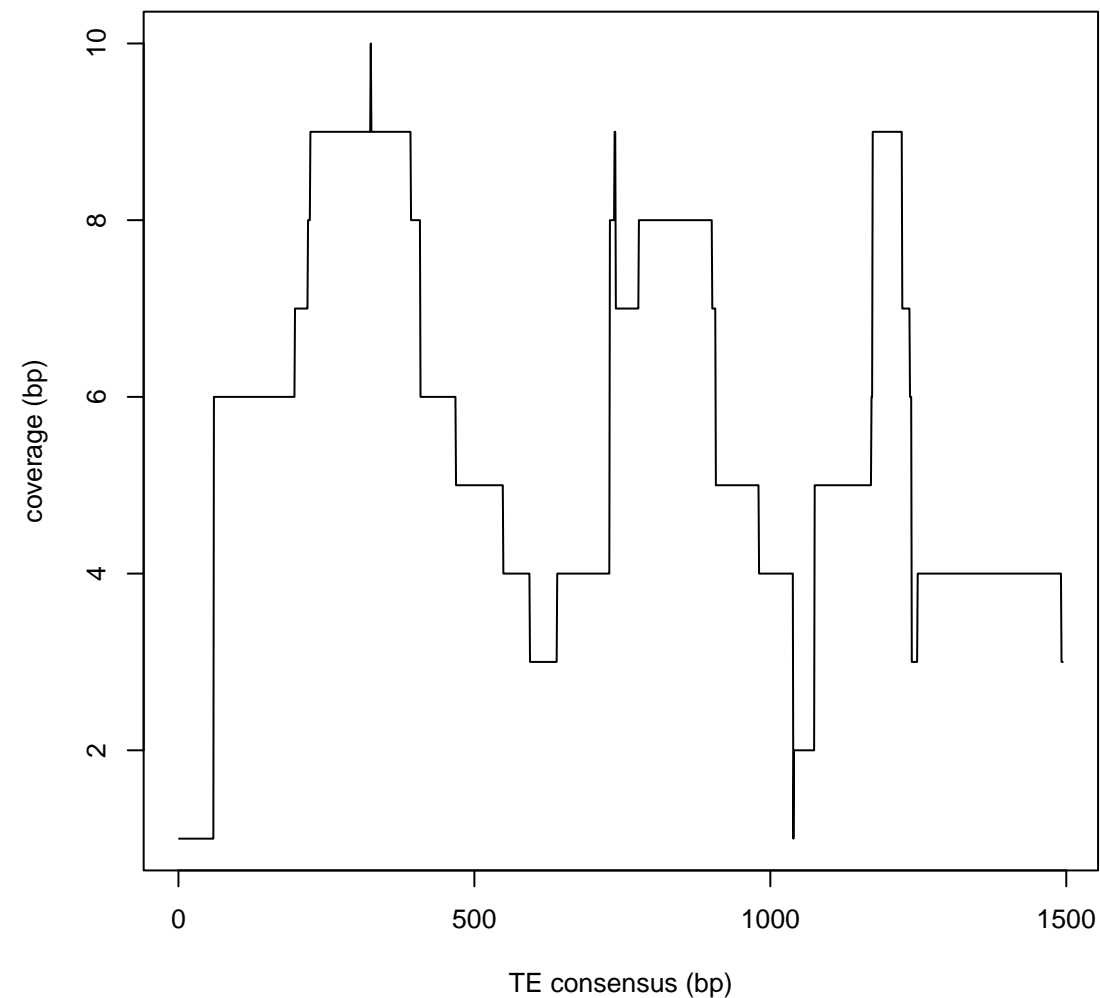


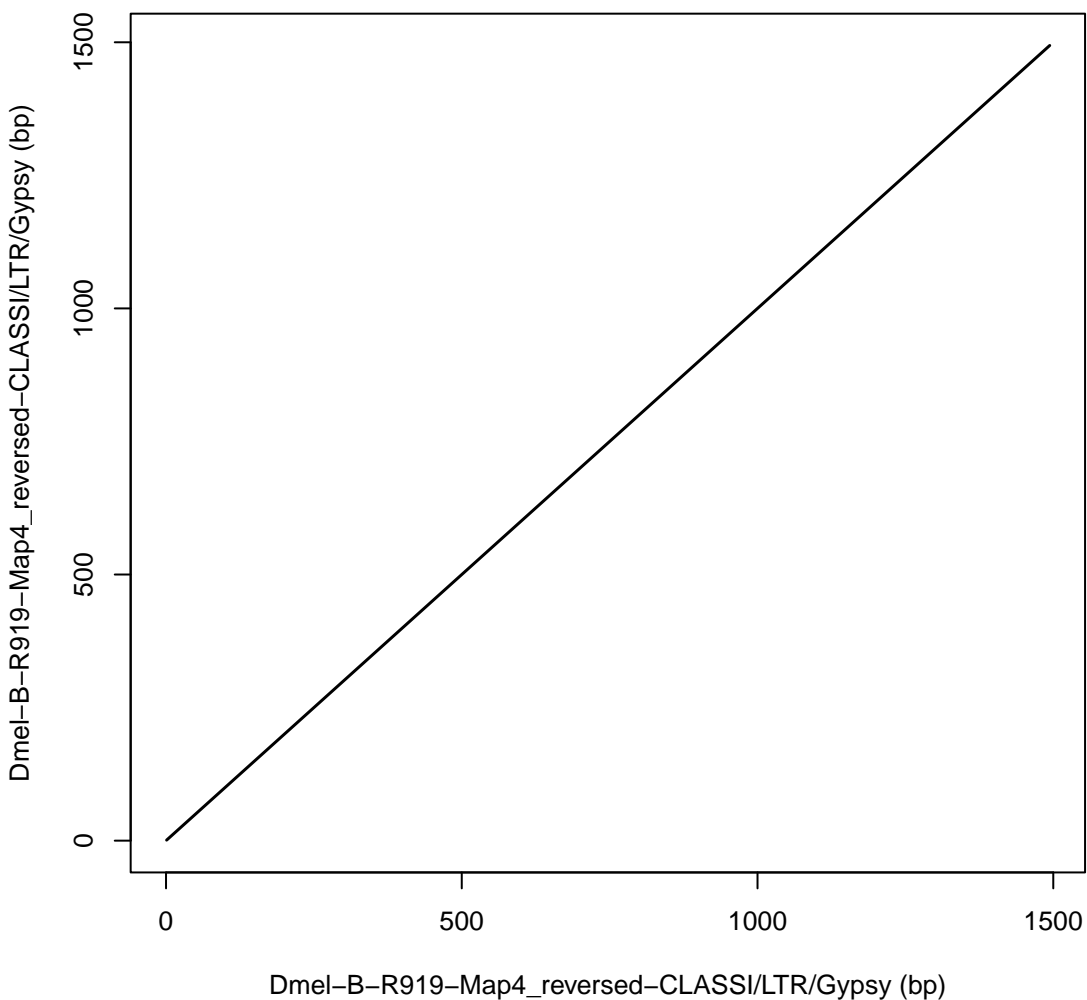
TE: Dmel-B-R919-Map4_reversed-CLASSI/LTR/Gypsy
consensus size: 1494bp; fragments: 26; full length: 1 (≥ 1344.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

