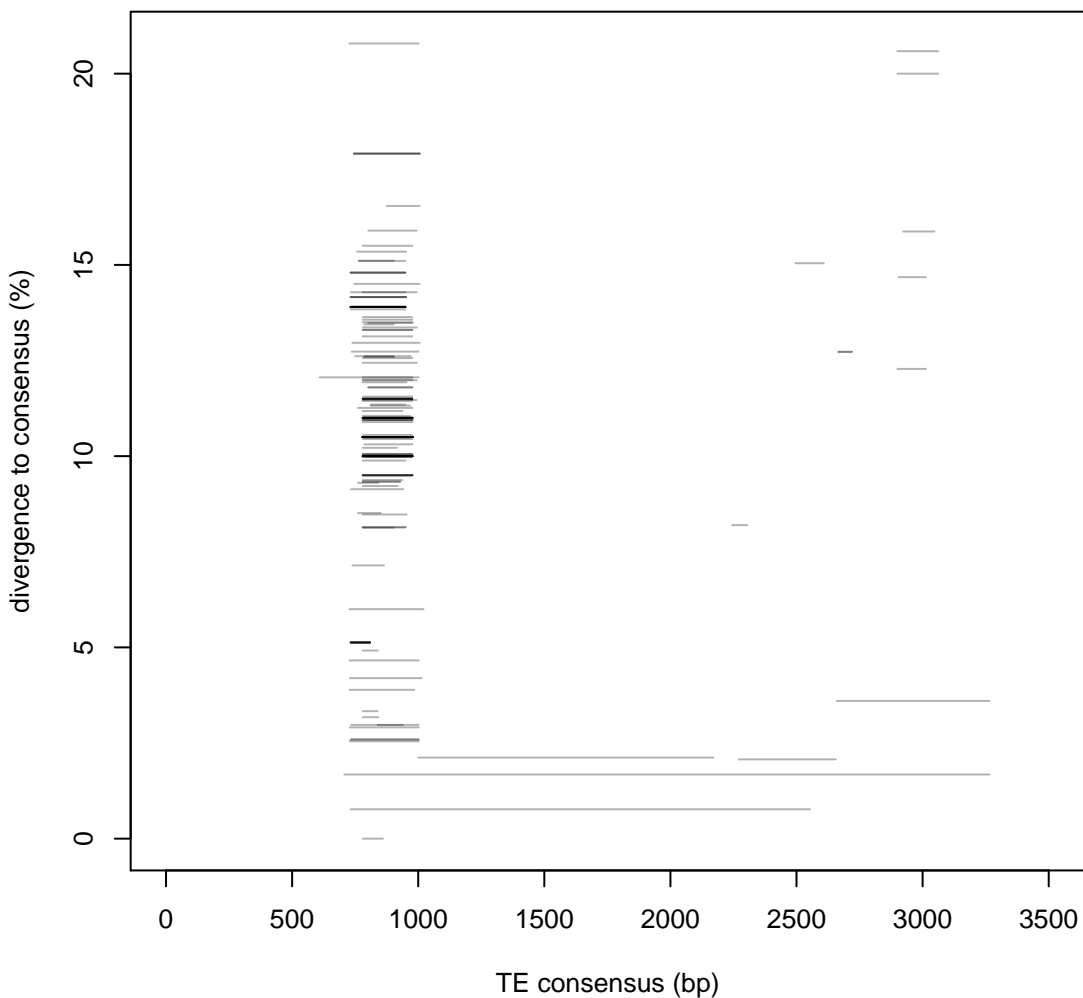
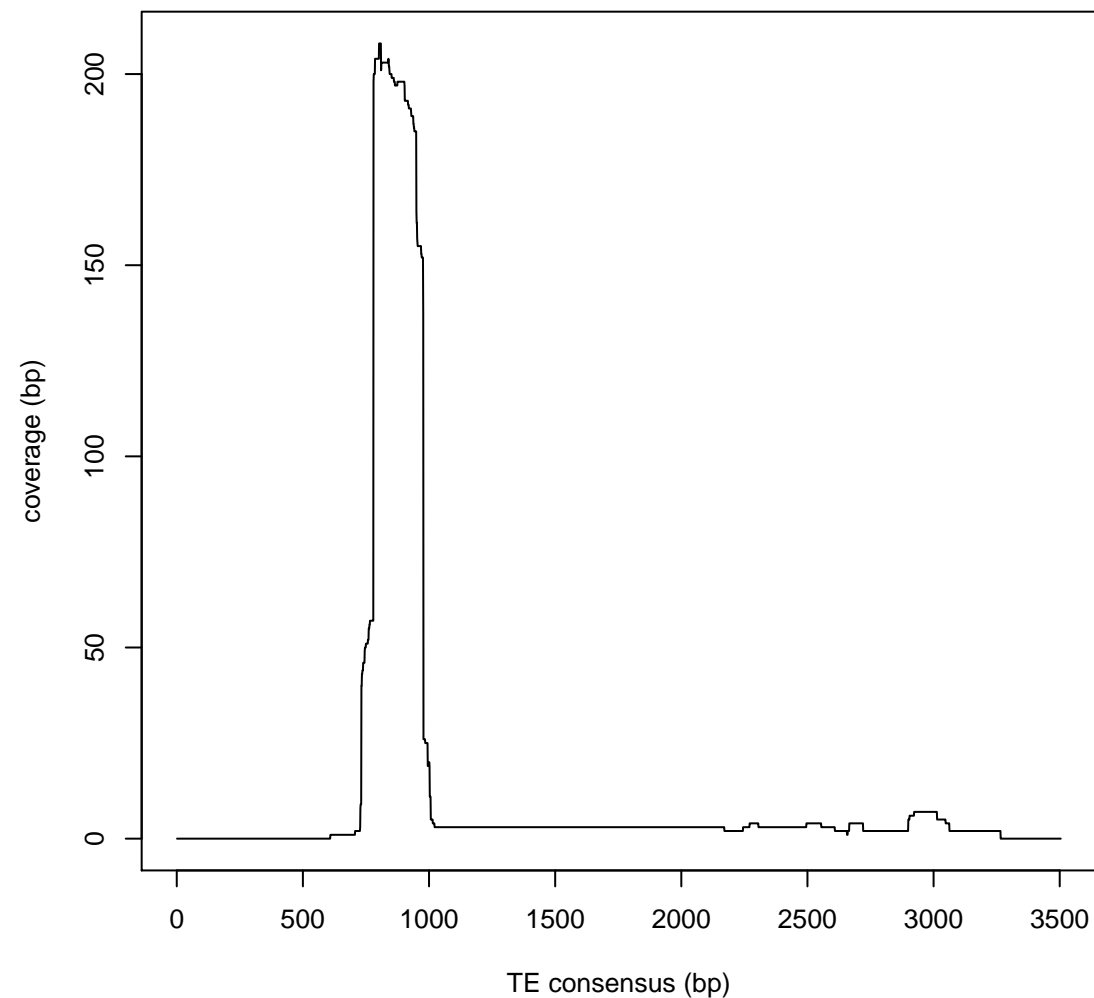


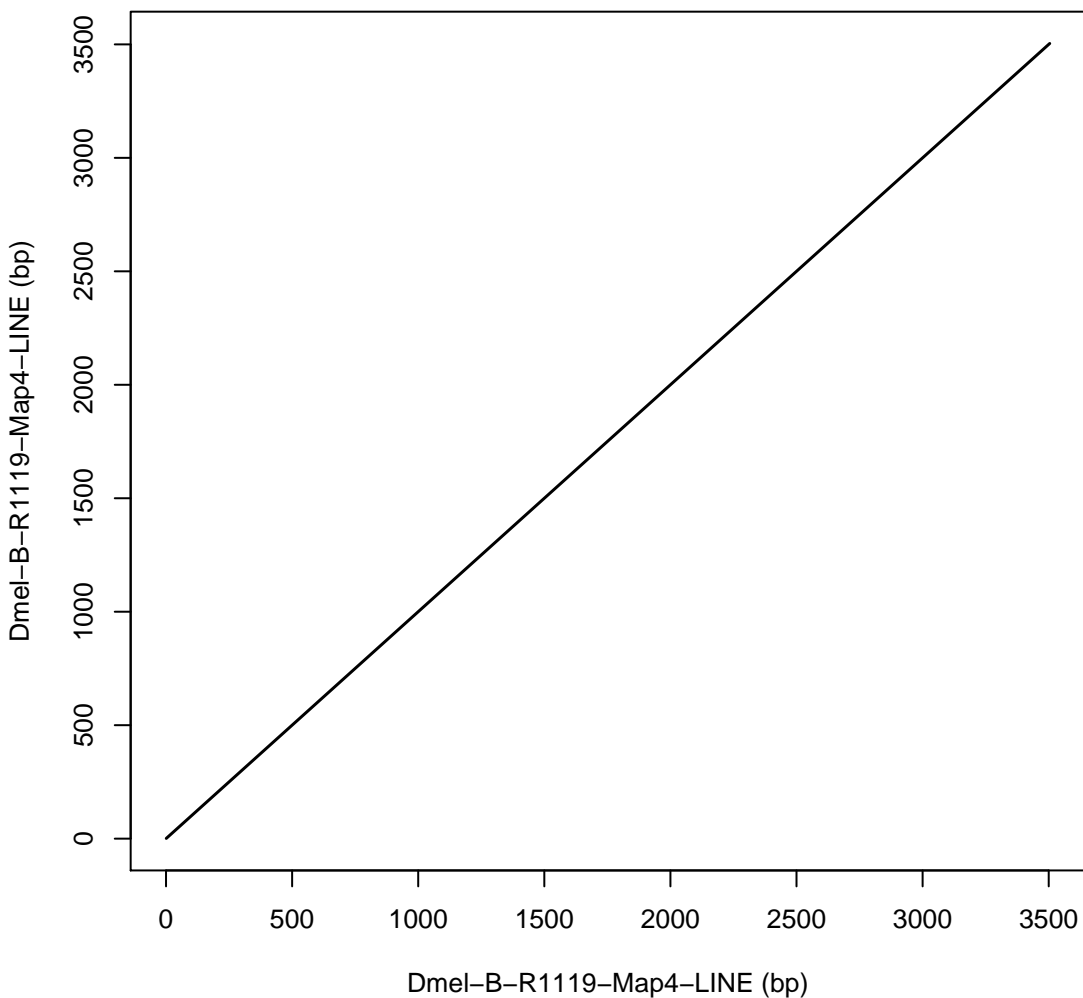
TE: Dmel-B-R1119-Map4-LINE
consensus size: 3504bp; fragments: 224; full length: 0 (≥ 3153.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

