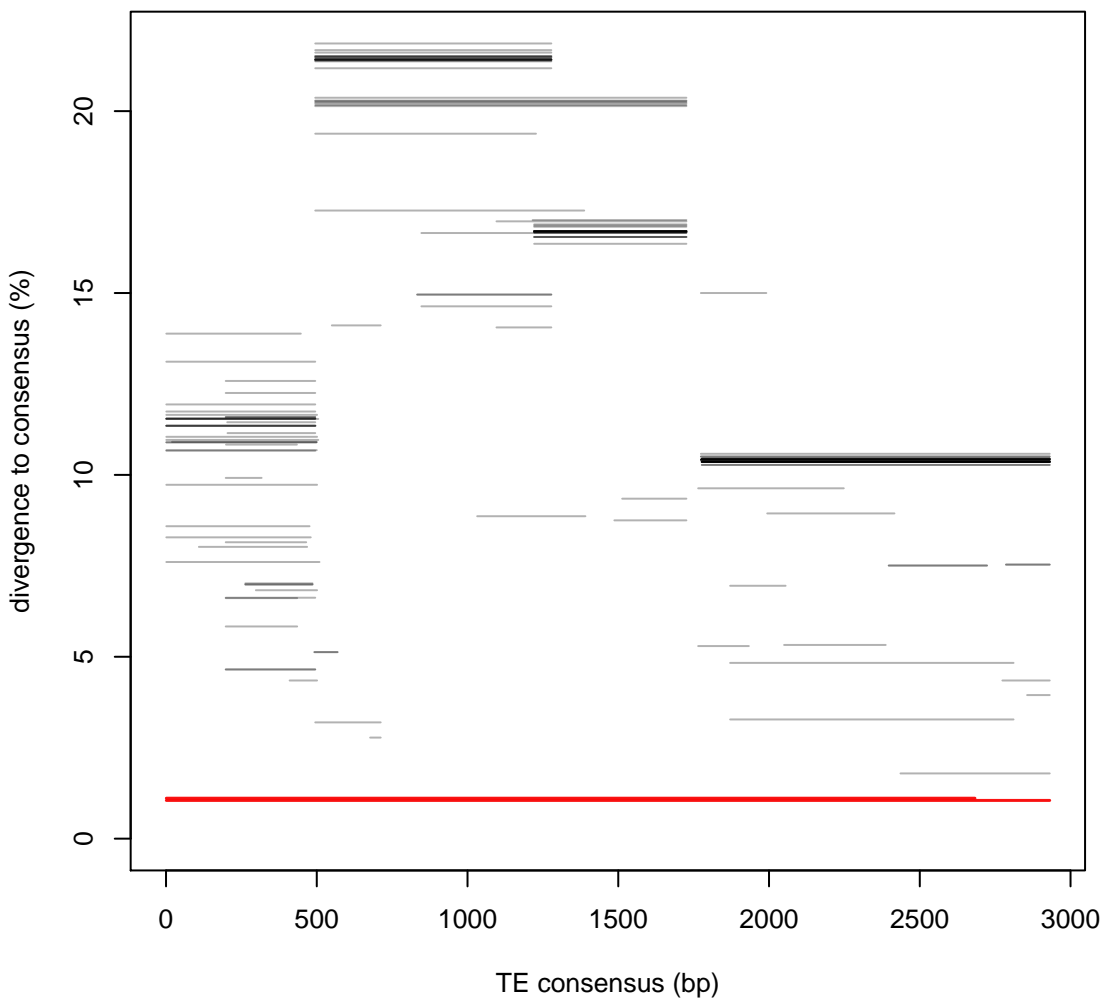
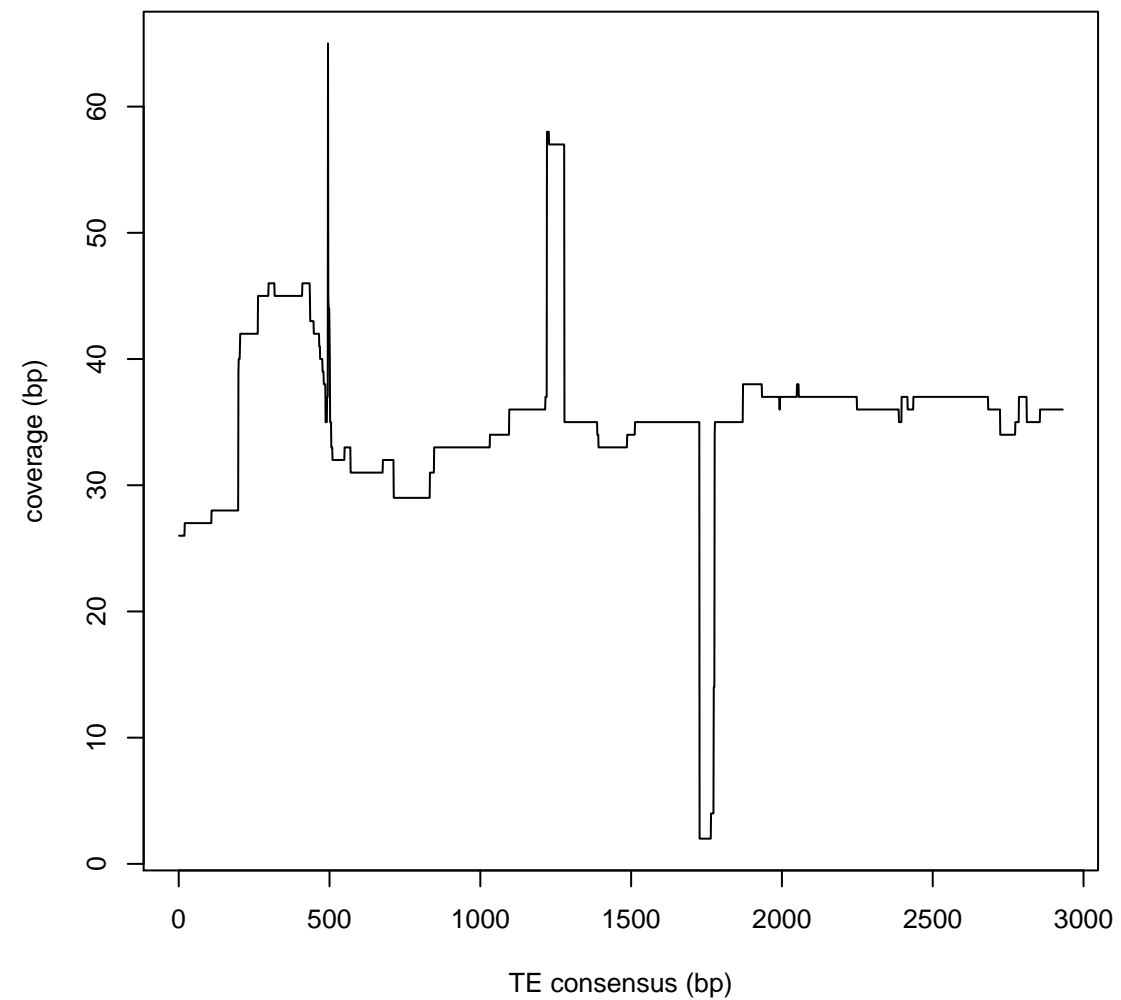


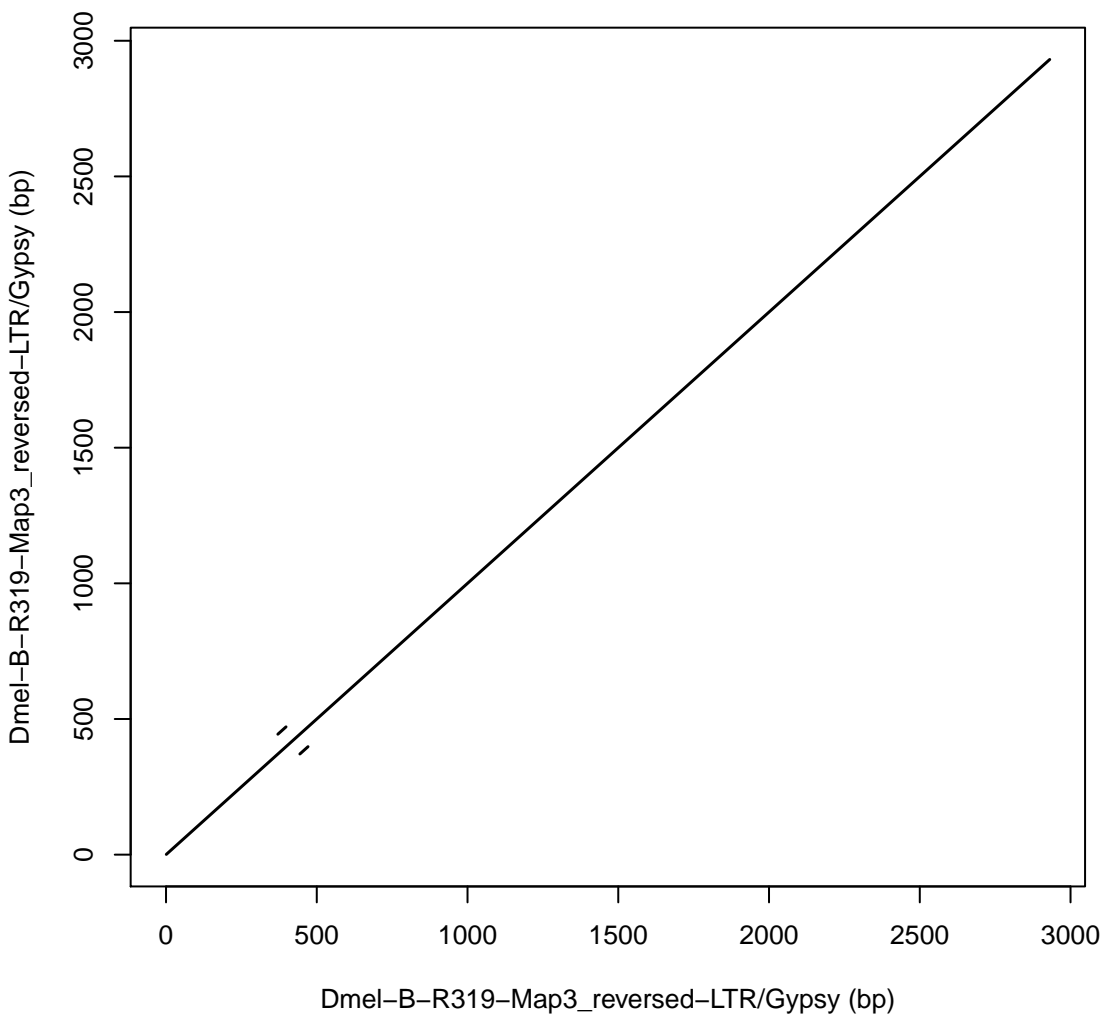
TE: Dmel-B-R319-Map3_reversed-LTR/Gypsy
consensus size: 2931bp; fragments: 155; full length: 2 (≥ 2637.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

