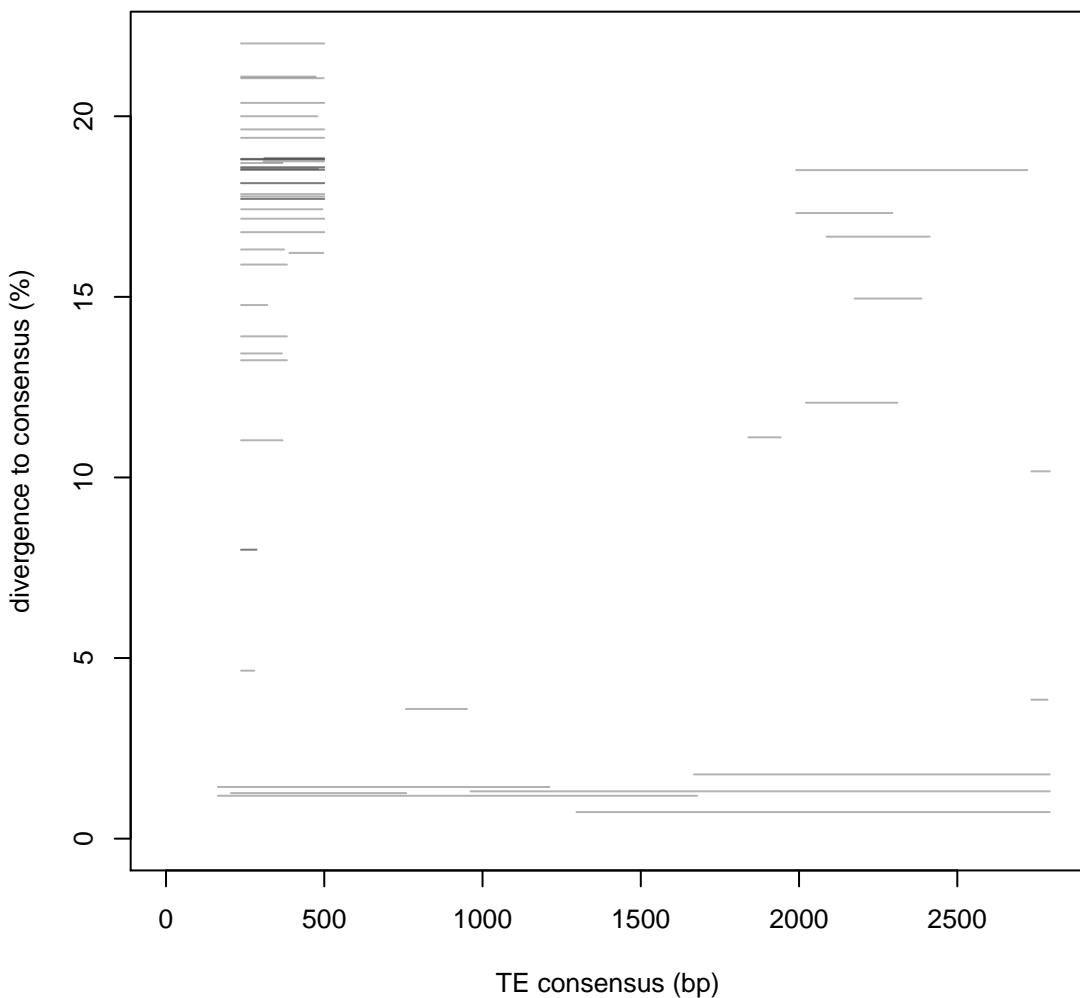
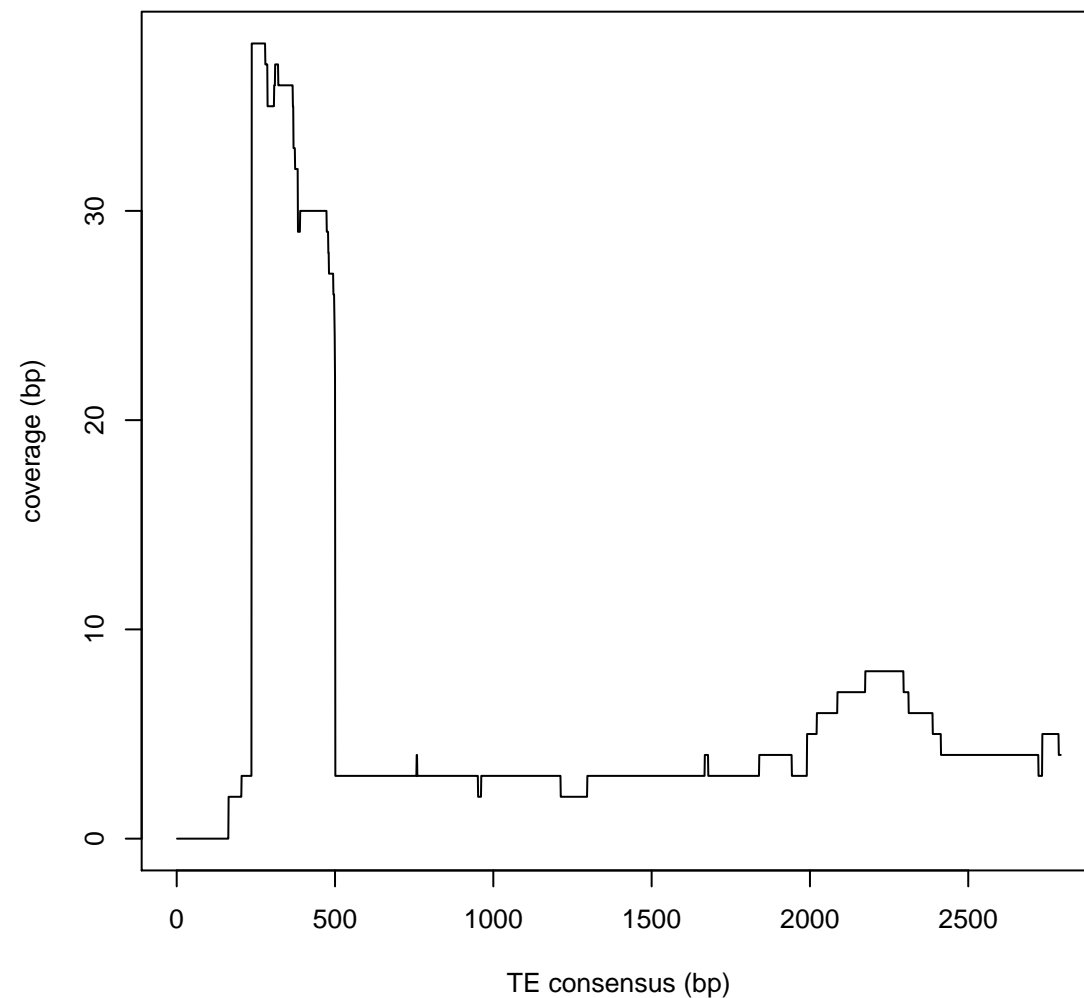


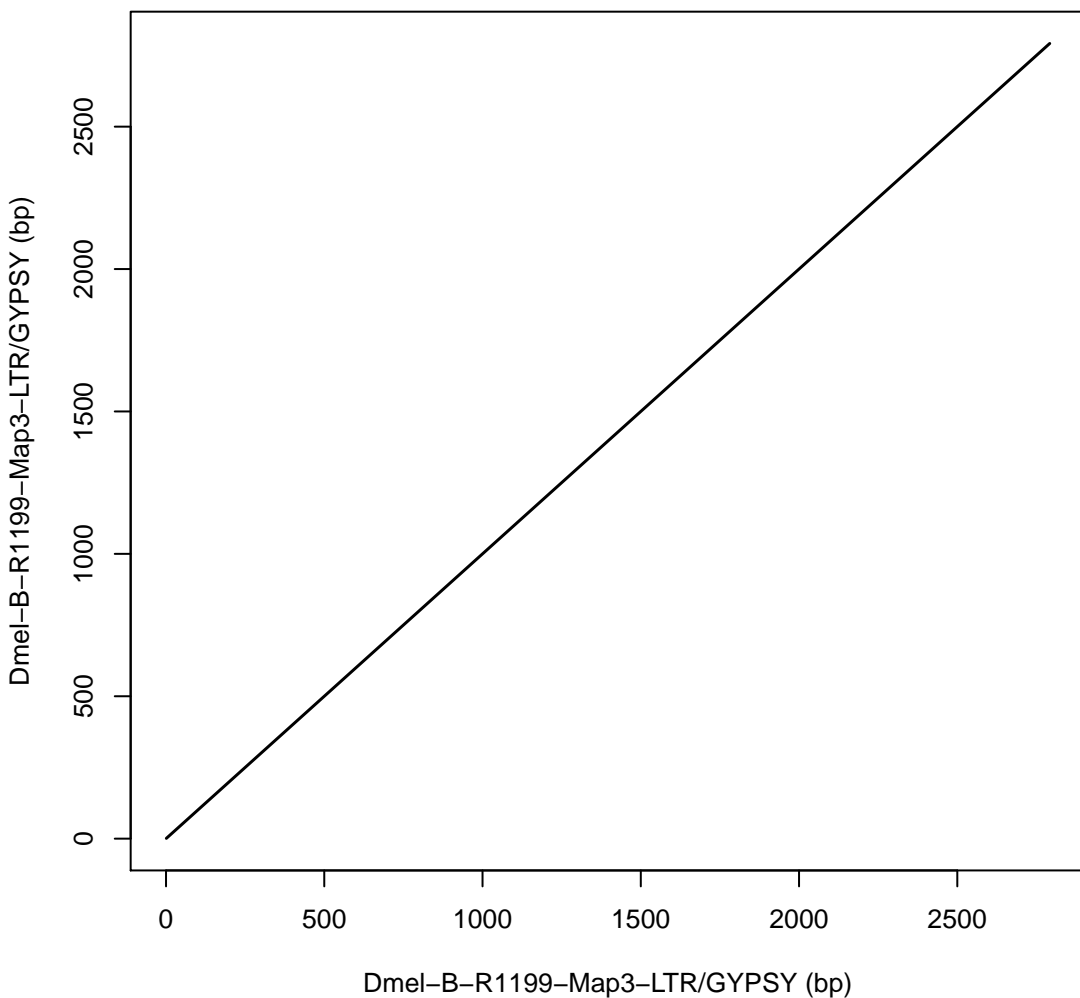
TE: Dmel-B-R1199-Map3-LTR/Gypsy
consensus size: 2792bp; fragments: 53; full length: 0 (>=2512.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

