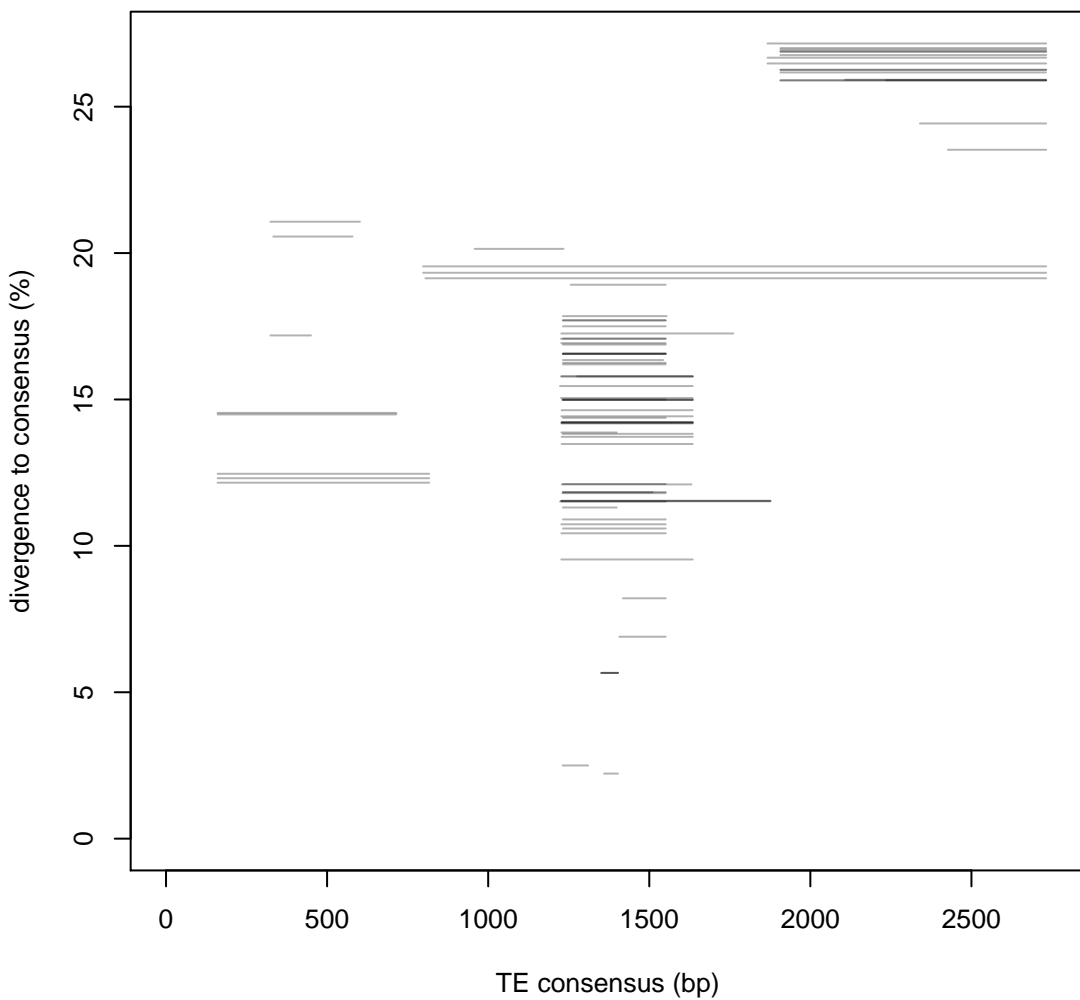
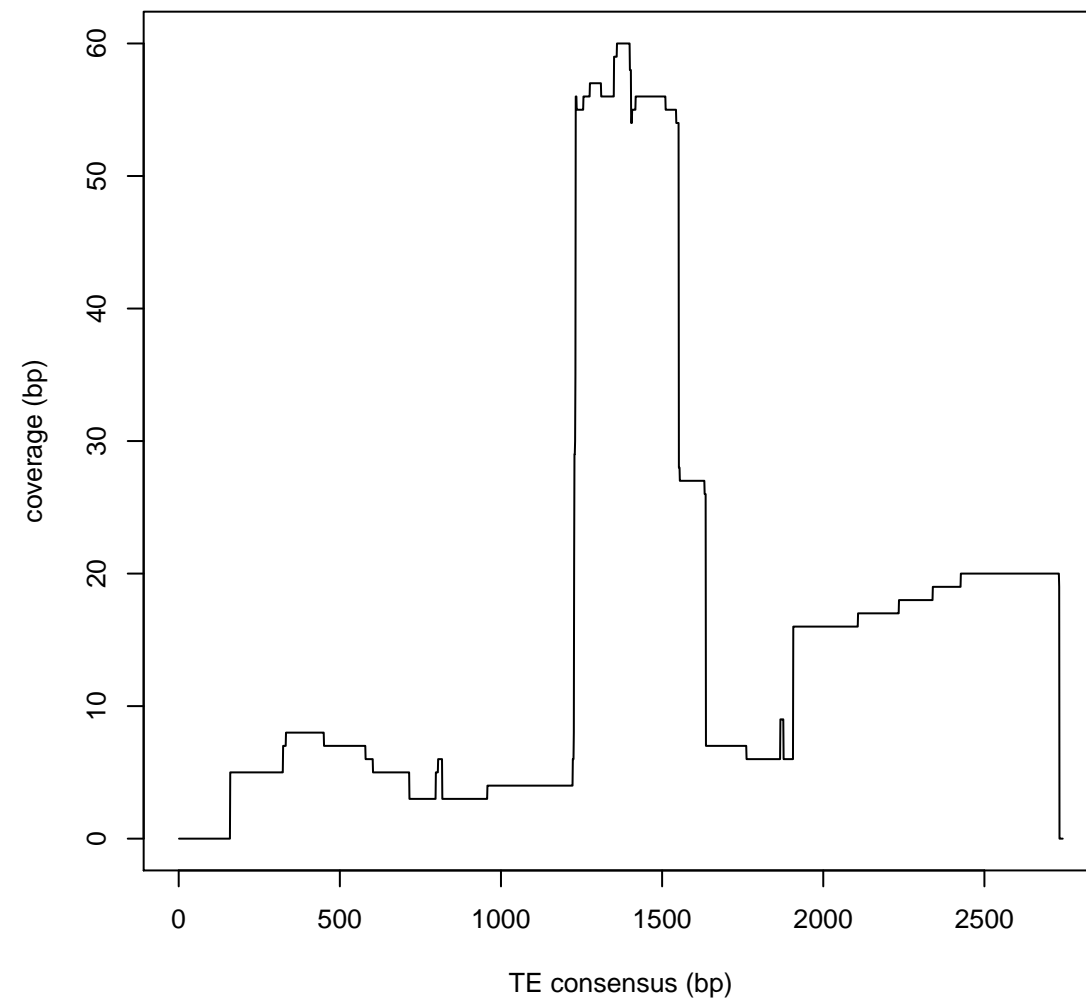


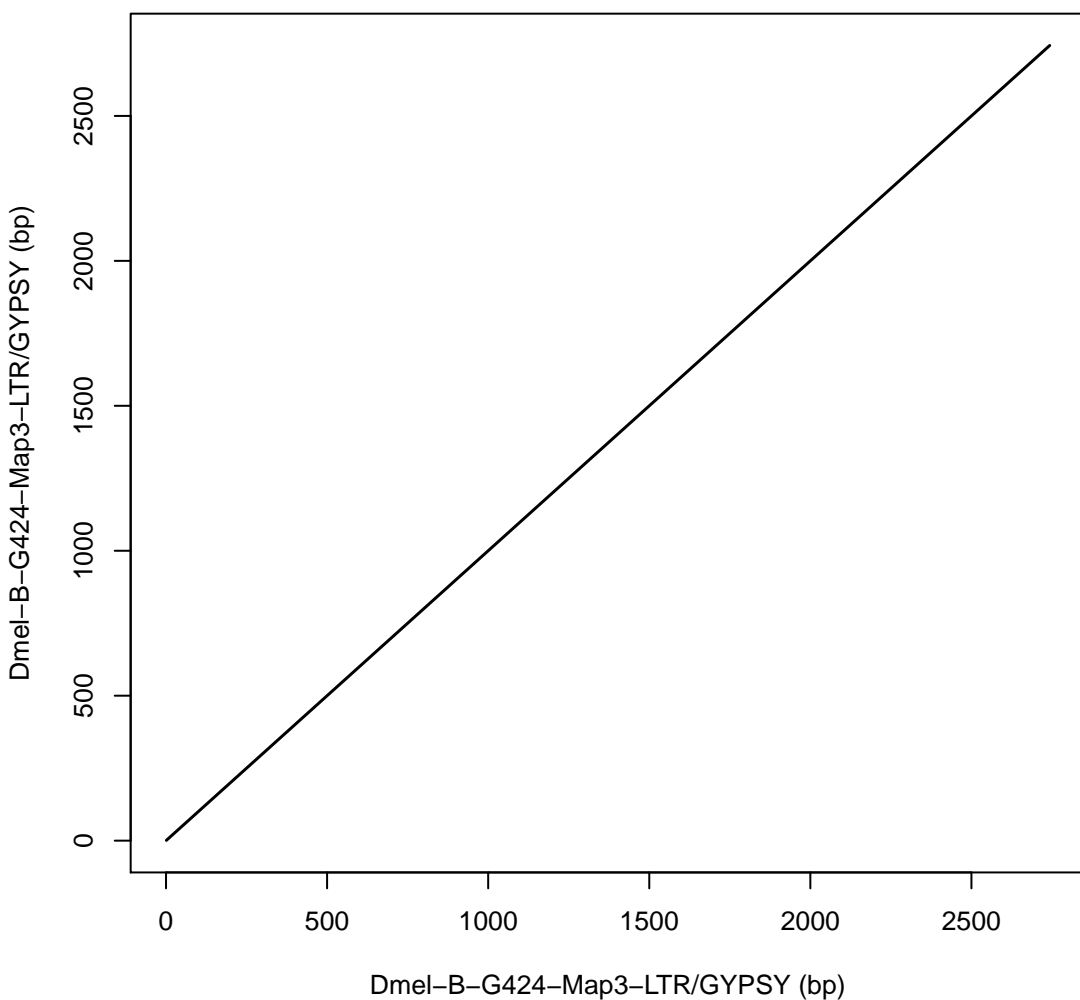
TE: Dmel-B-G424-Map3-LTR/GYPSY
consensus size: 2743bp; fragments: 89; full length: 0 (≥ 2468.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

