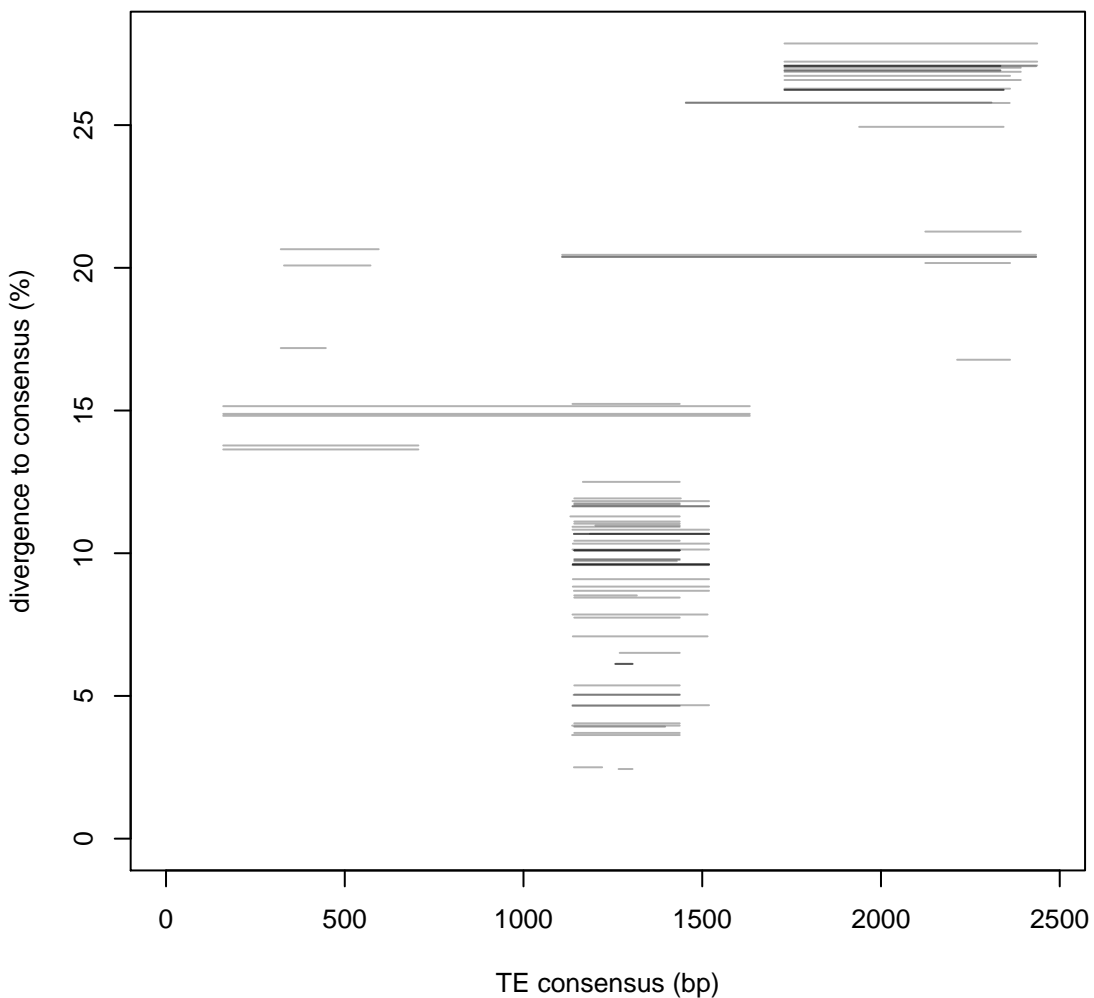
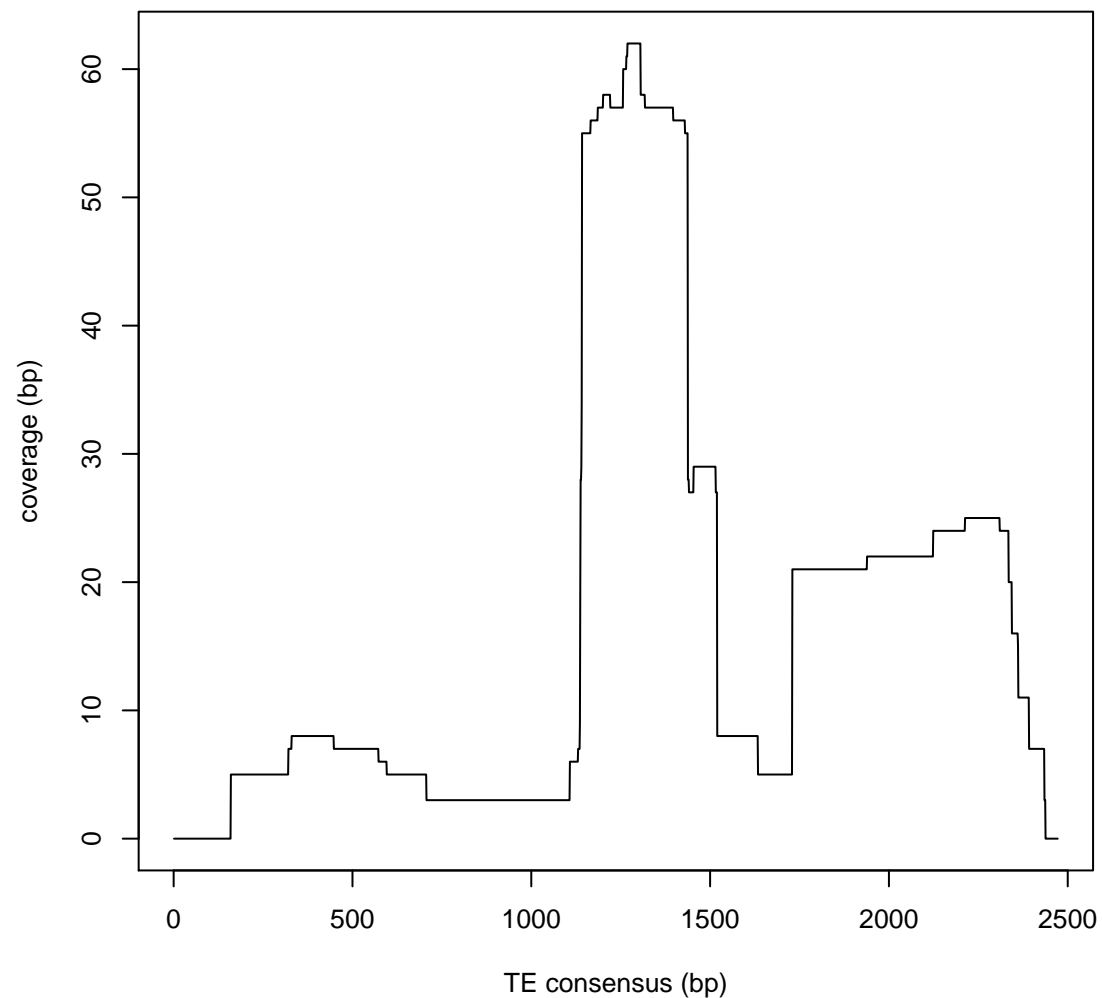


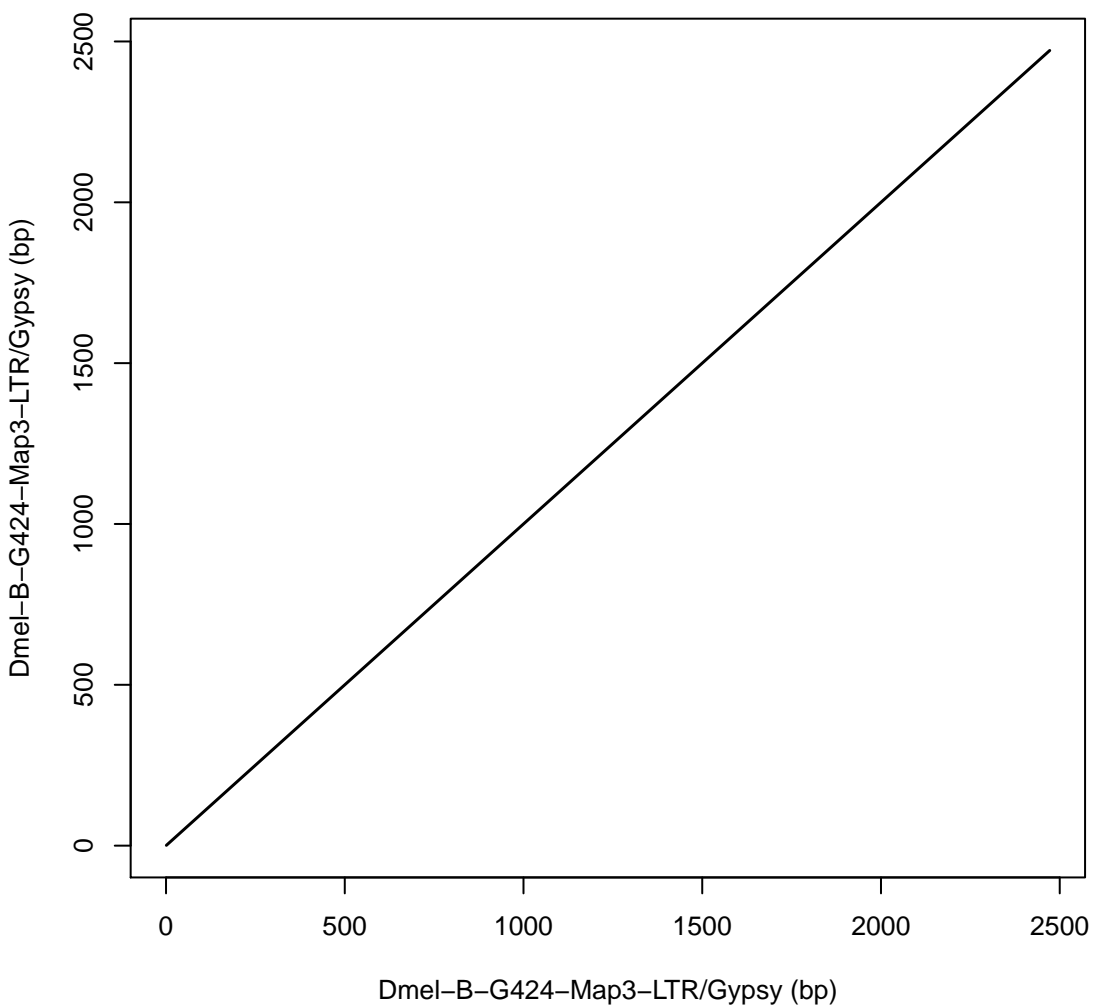
TE: Dmel-B-G424-Map3-LTR/Gypsy
consensus size: 2472bp; fragments: 90; full length: 0 (≥ 2224.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

