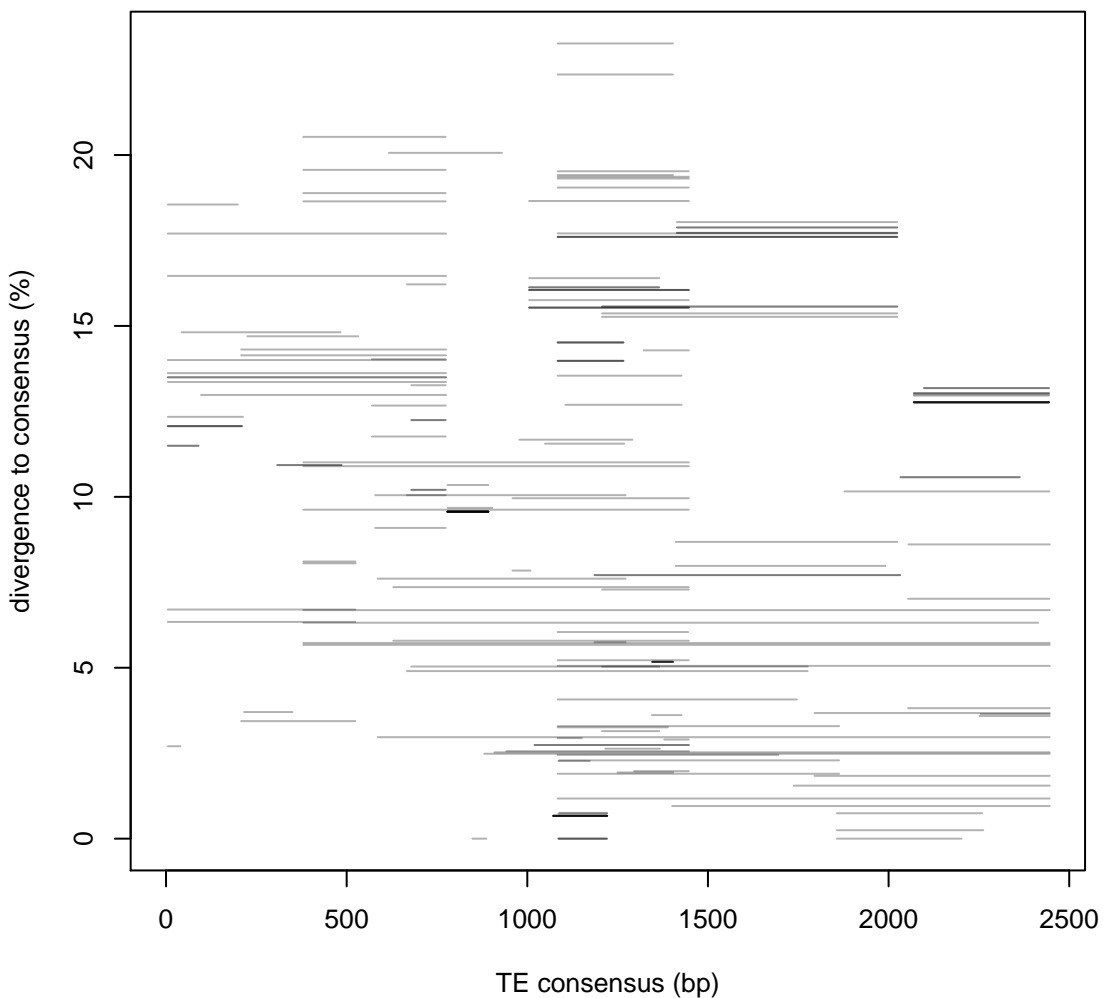
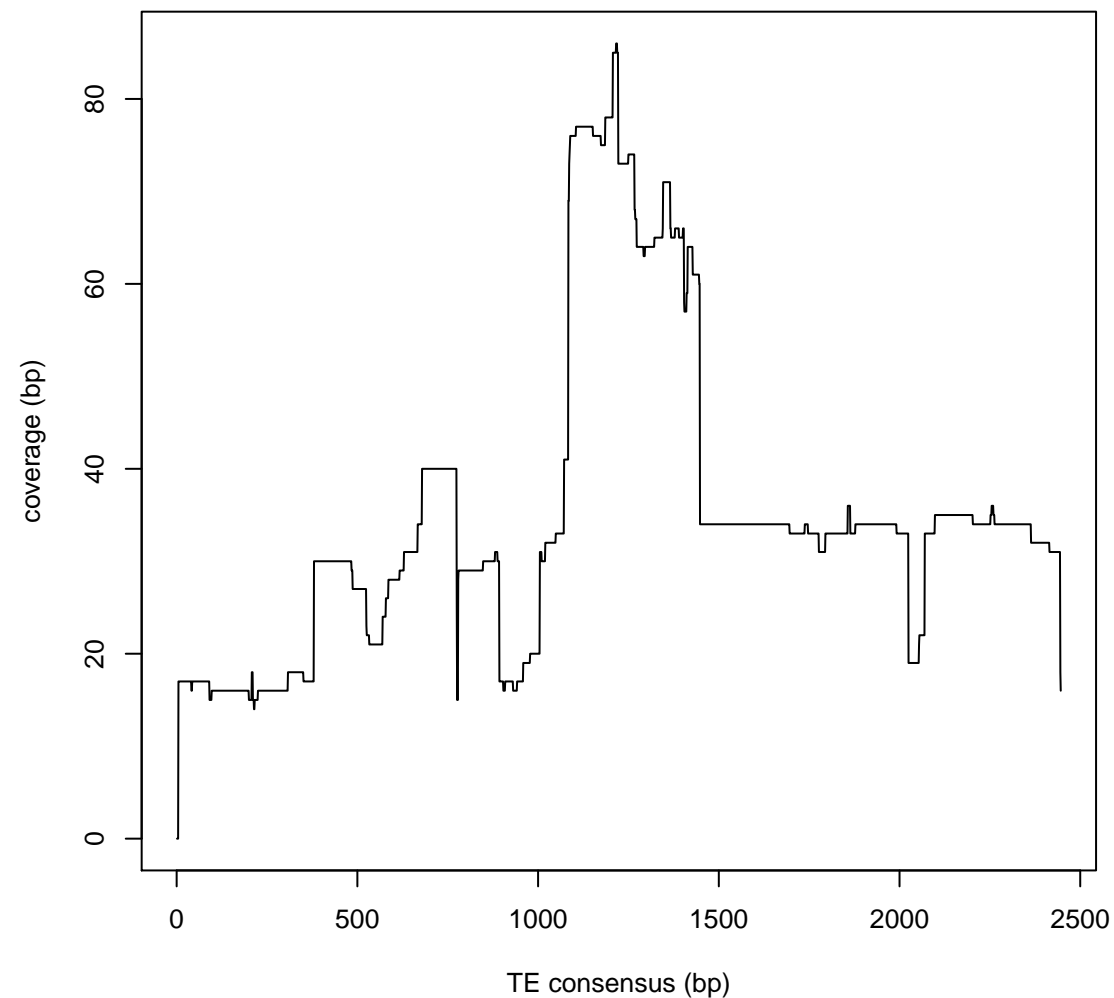


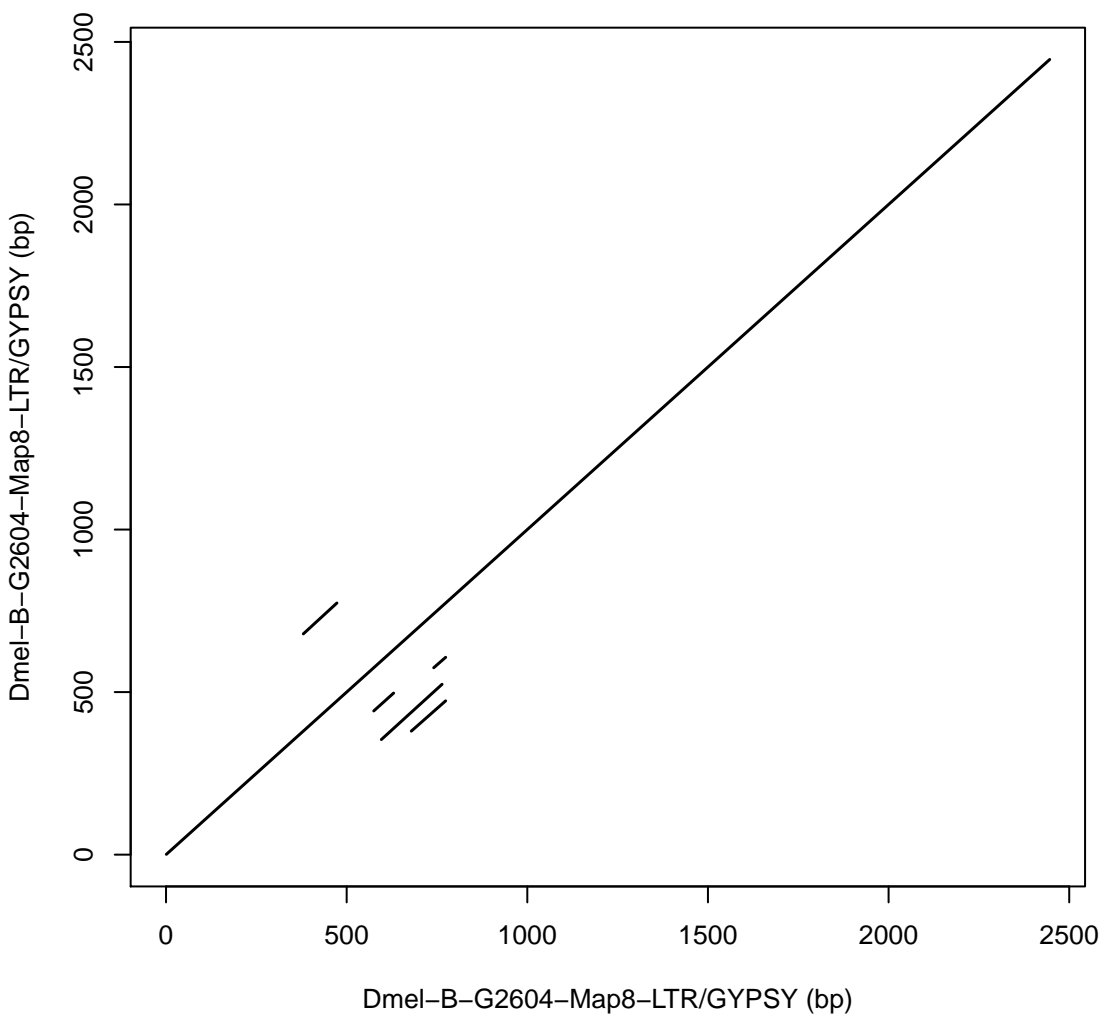
TE: Dmel-B-G2604-Map8-LTR/GYPSY
consensus size: 2446bp; fragments: 193; full length: 0 (>=2201.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

