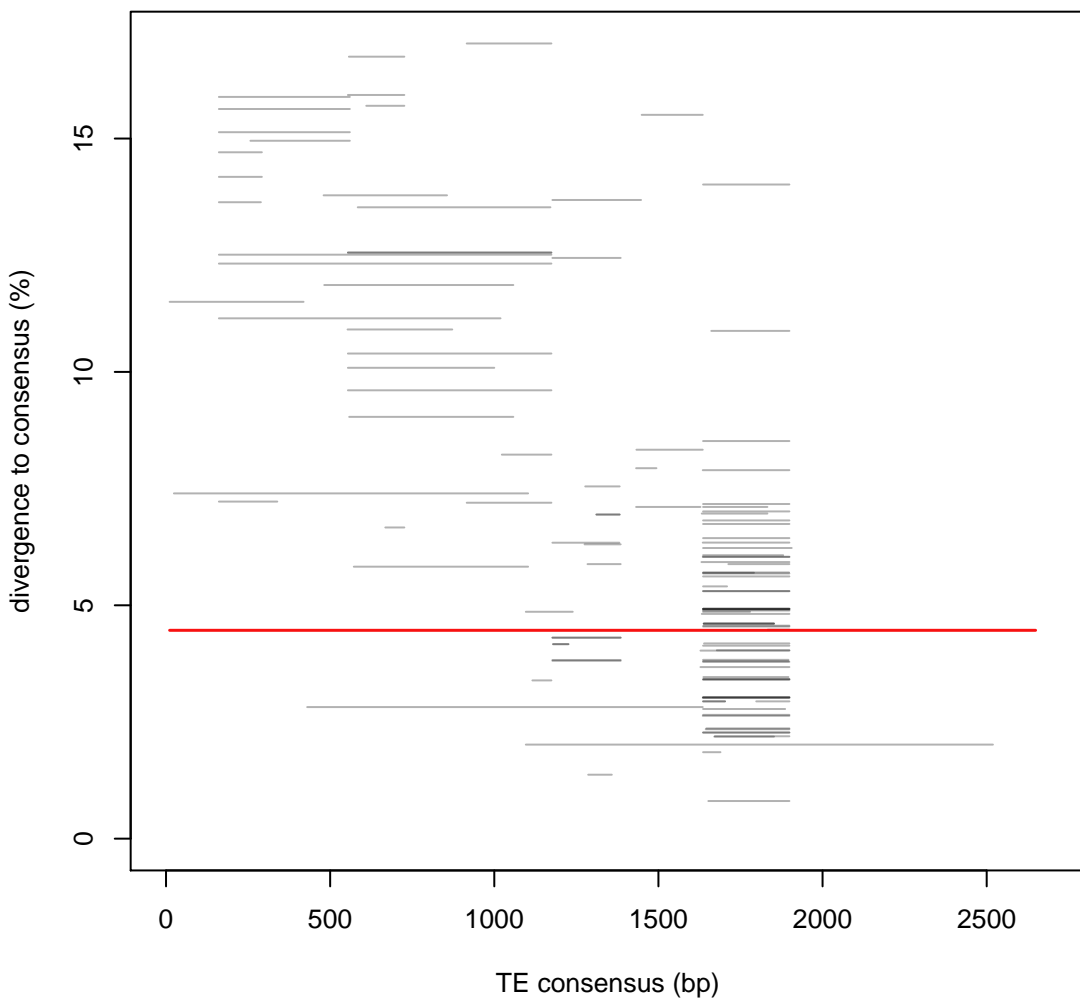
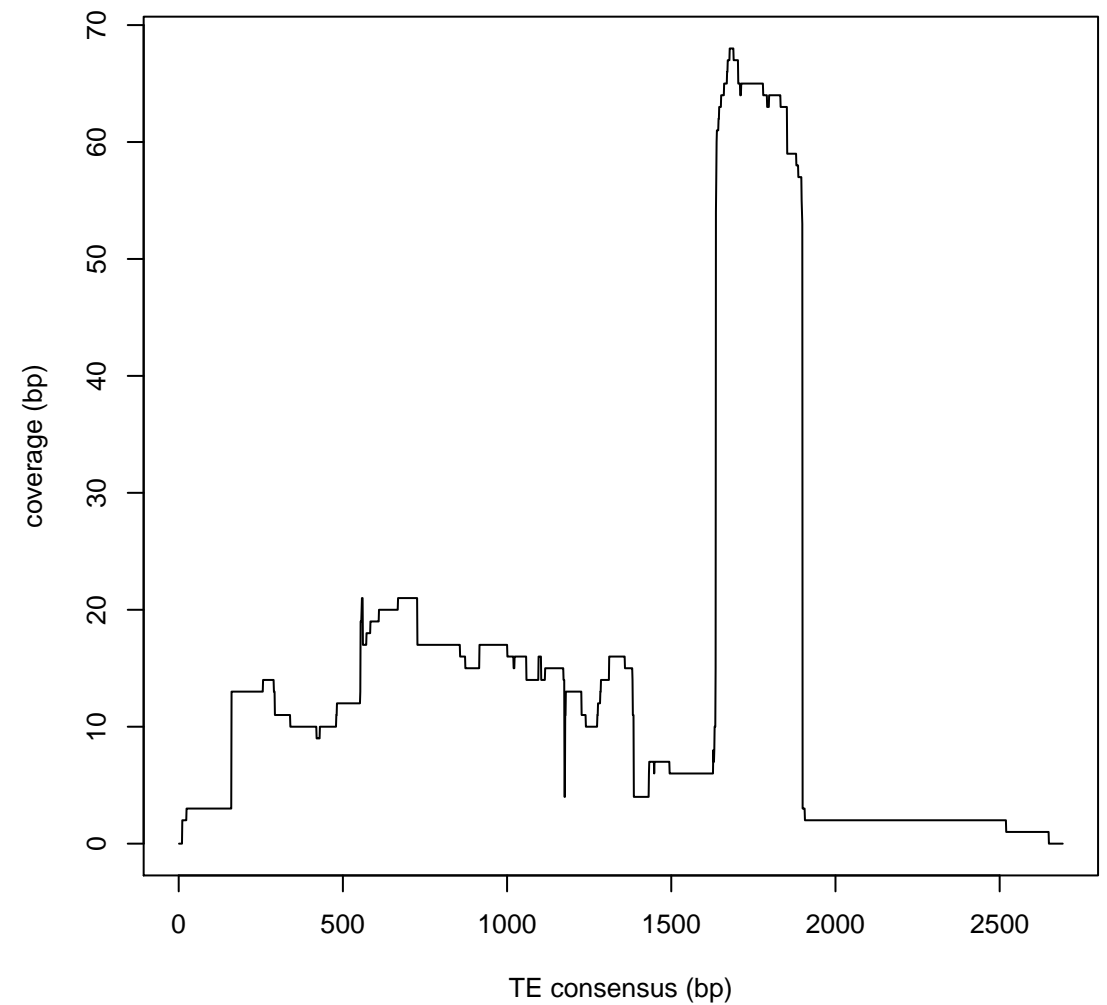


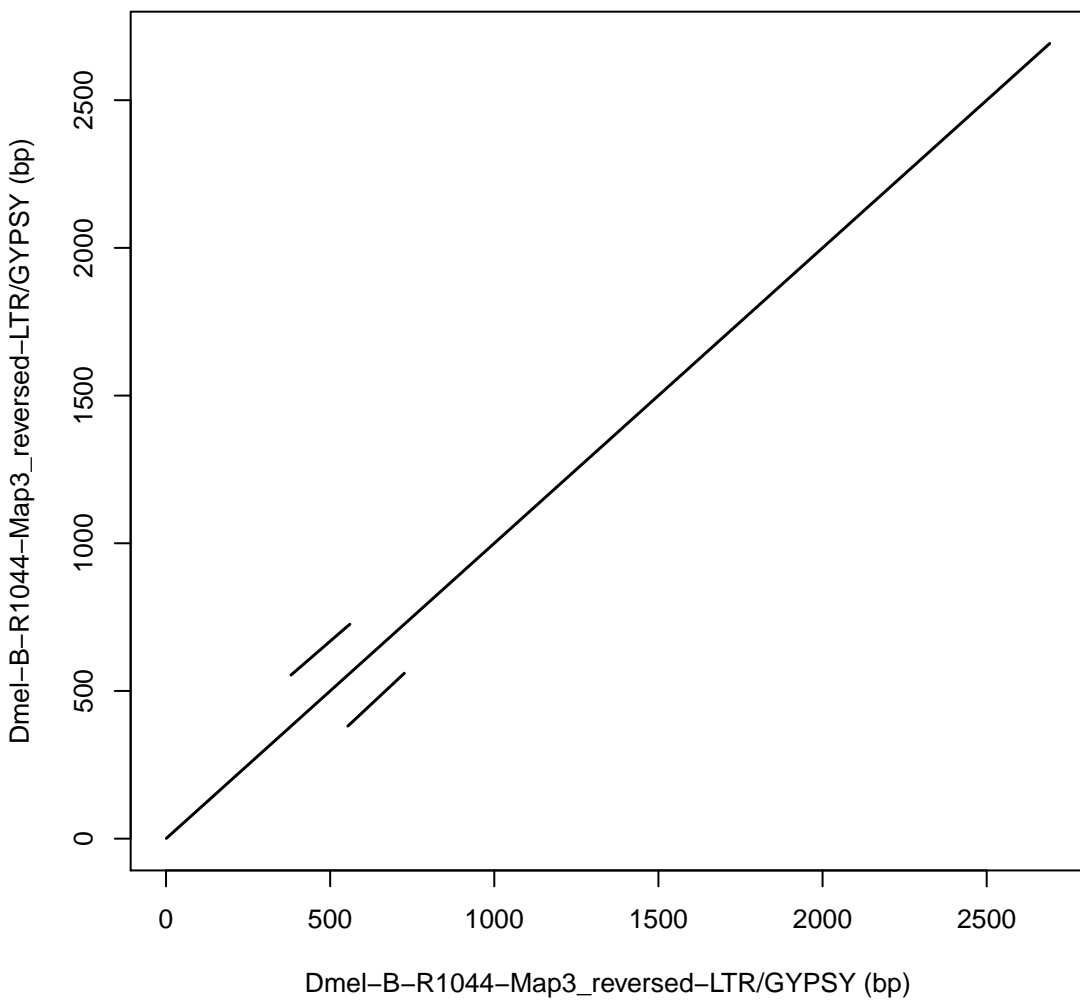
TE: Dmel-B-R1044-Map3_reversed-LTR/GYPSY
consensus size: 2692bp; fragments: 124; full length: 1 (>=2422.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

