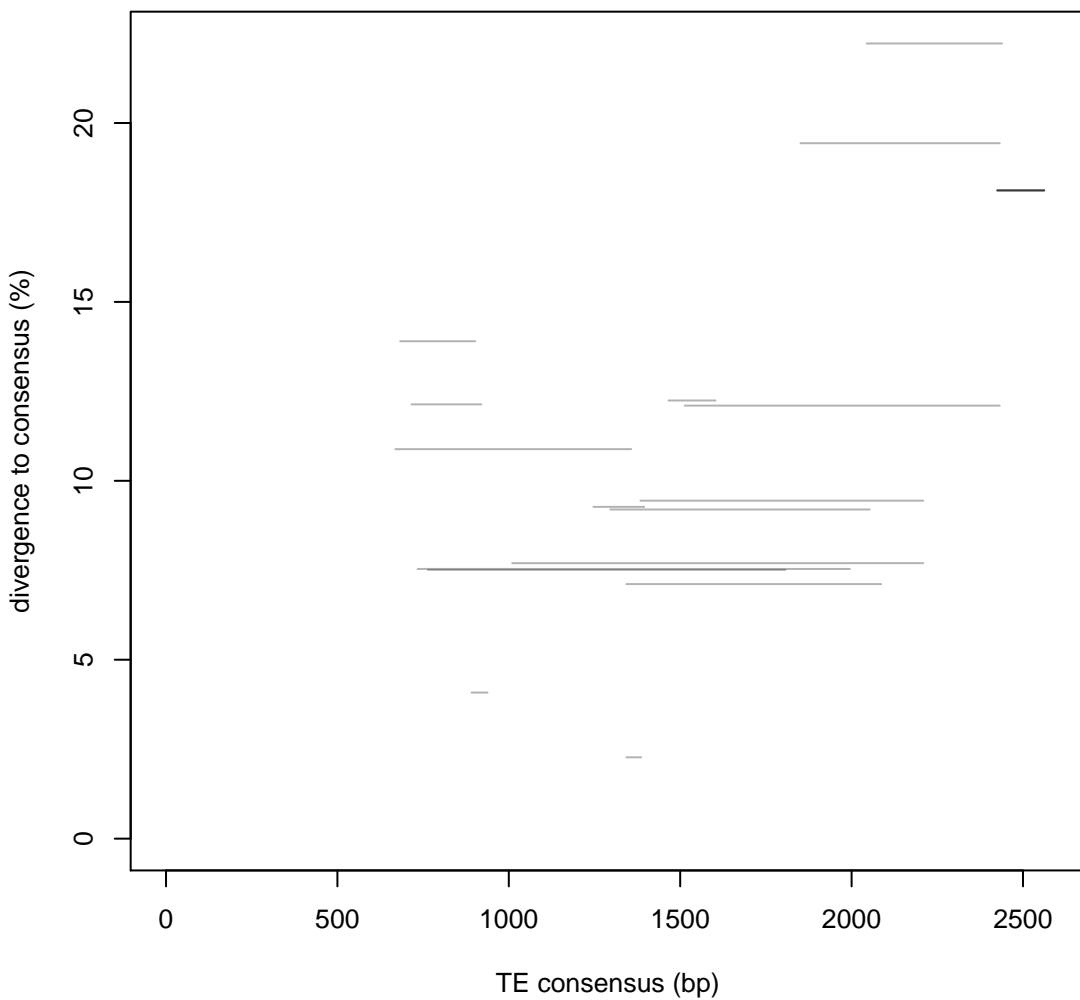
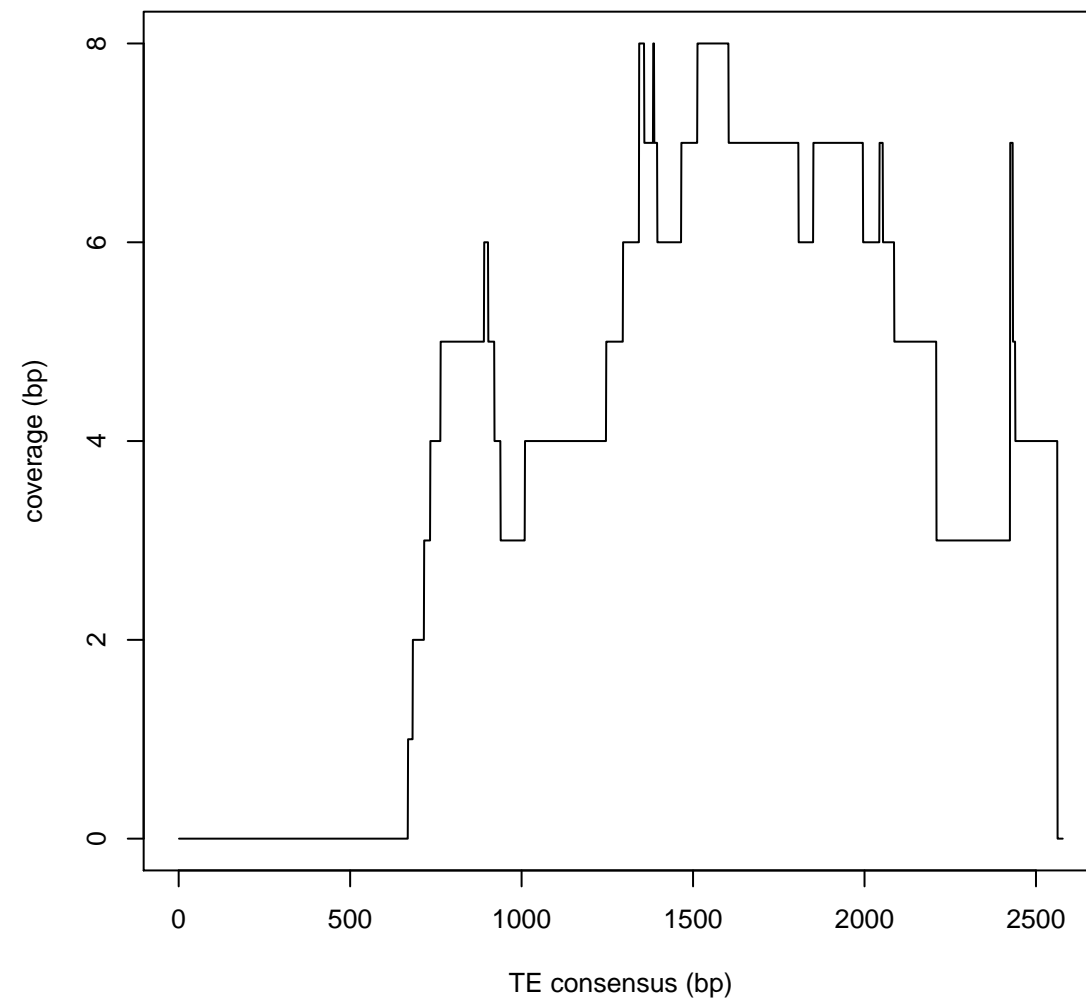


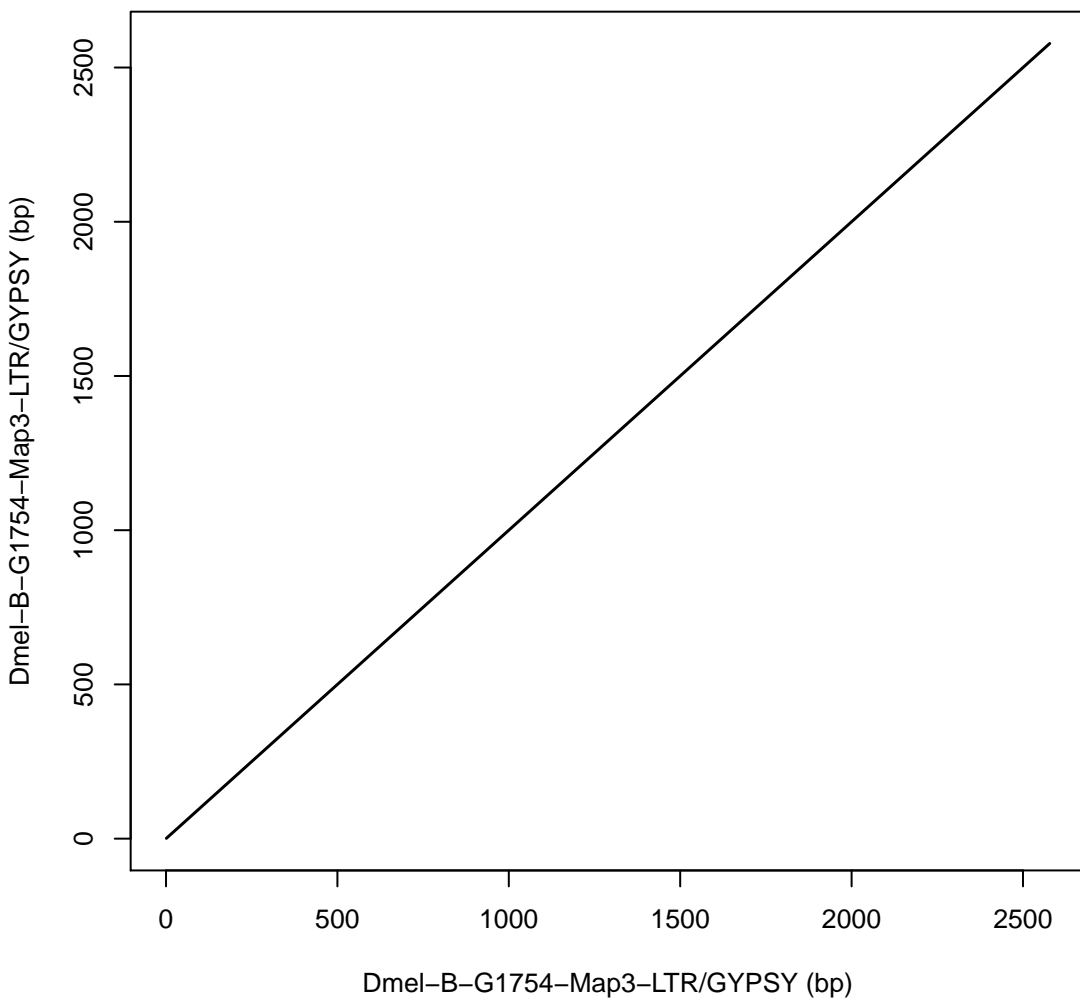
TE: Dmel-B-G1754-Map3-LTR/GYPSY
consensus size: 2578bp; fragments: 20; full length: 0 (>=2320.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

