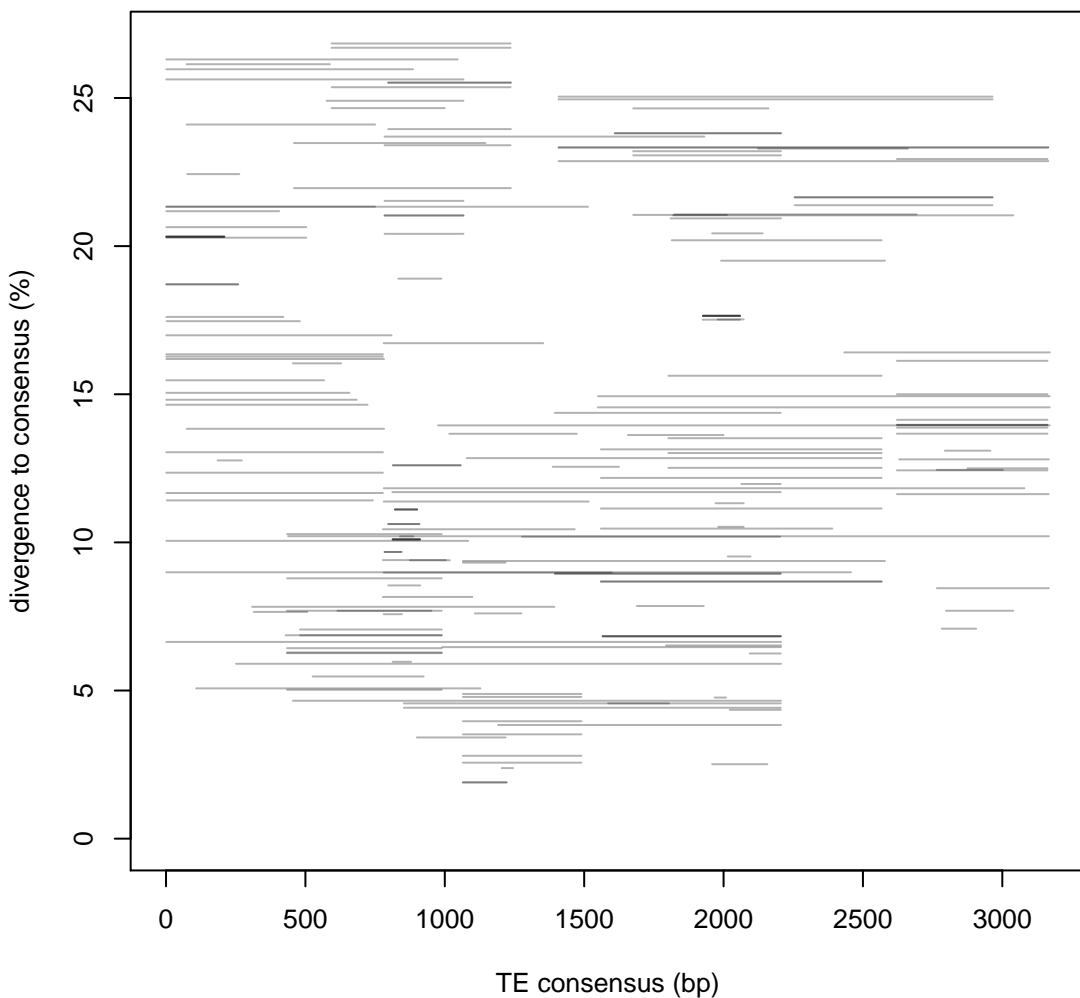
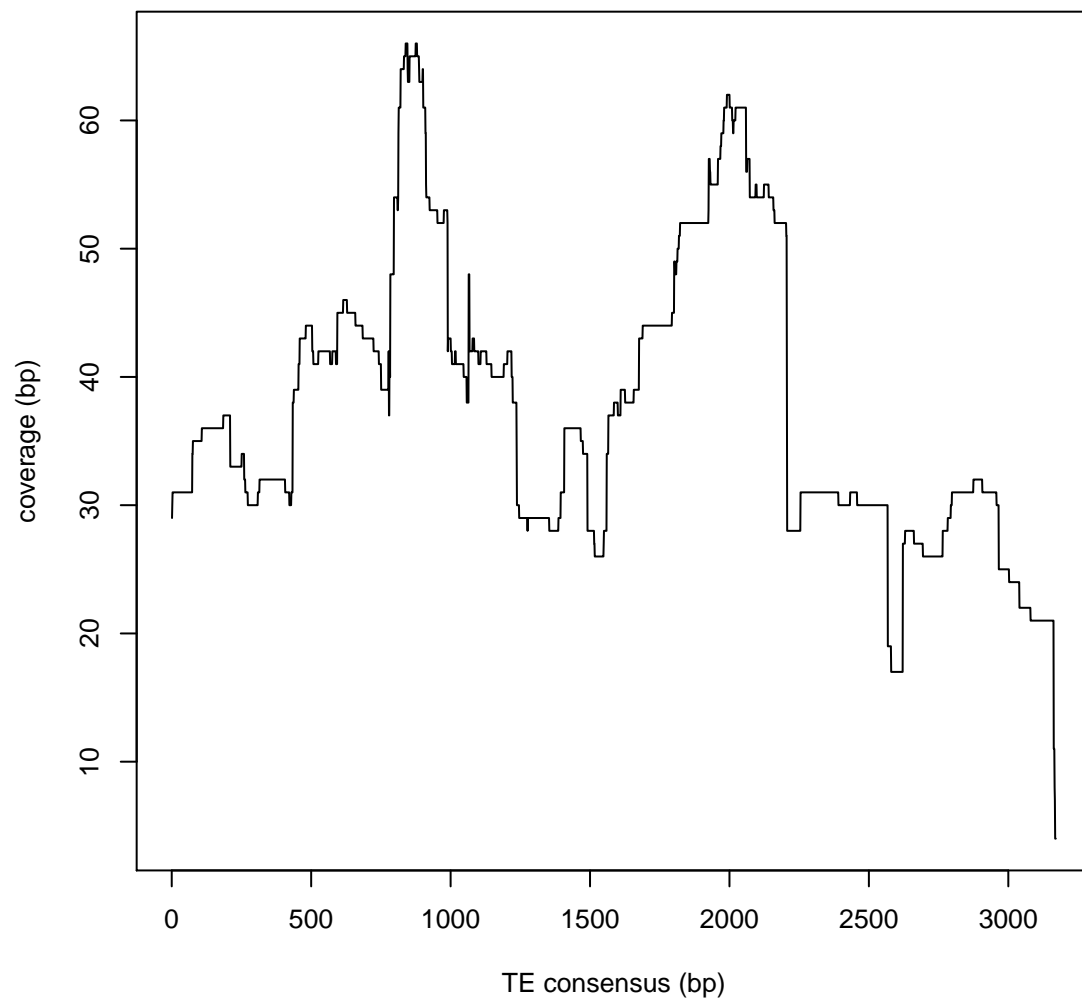


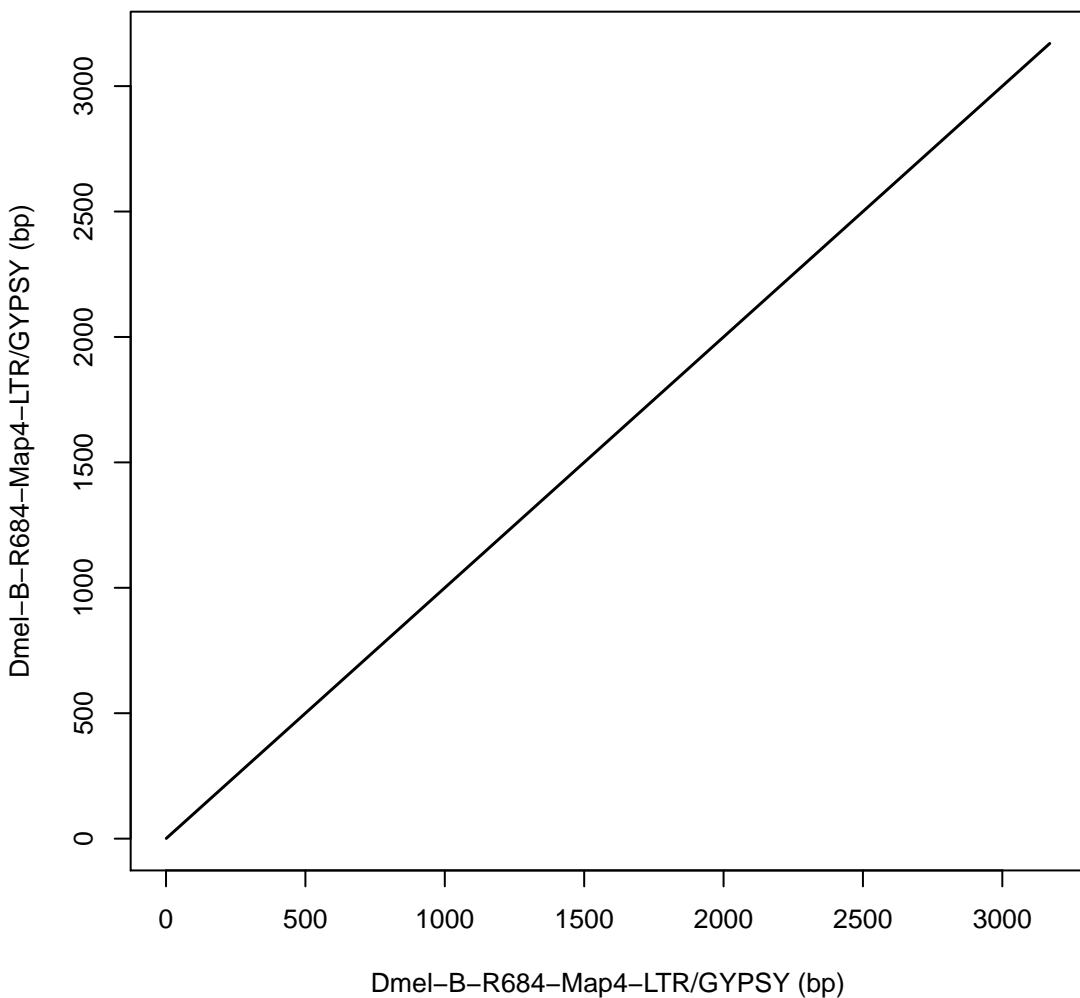
TE: Dmel-B-R684-Map4-LTR/GYPSY
consensus size: 3170bp; fragments: 195; full length: 0 (>=2853bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

