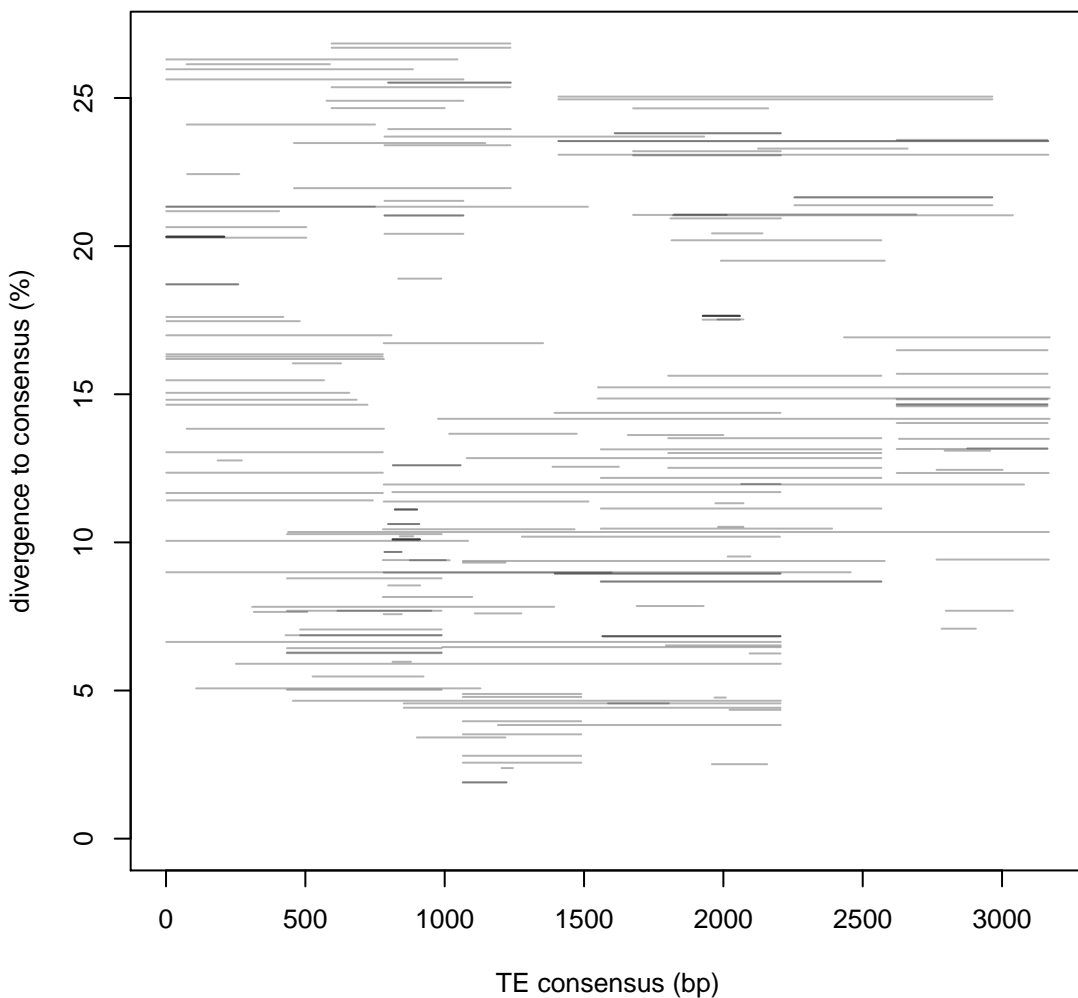
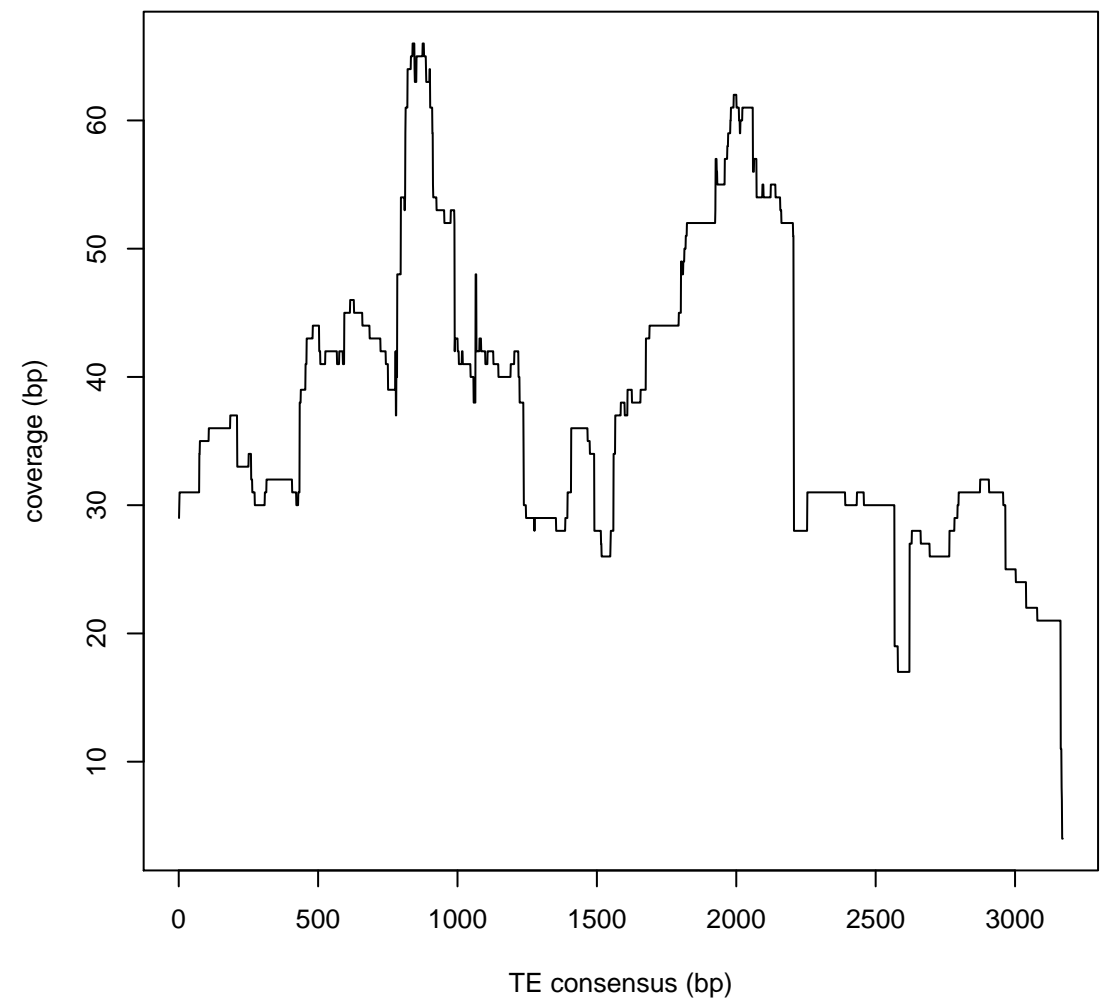


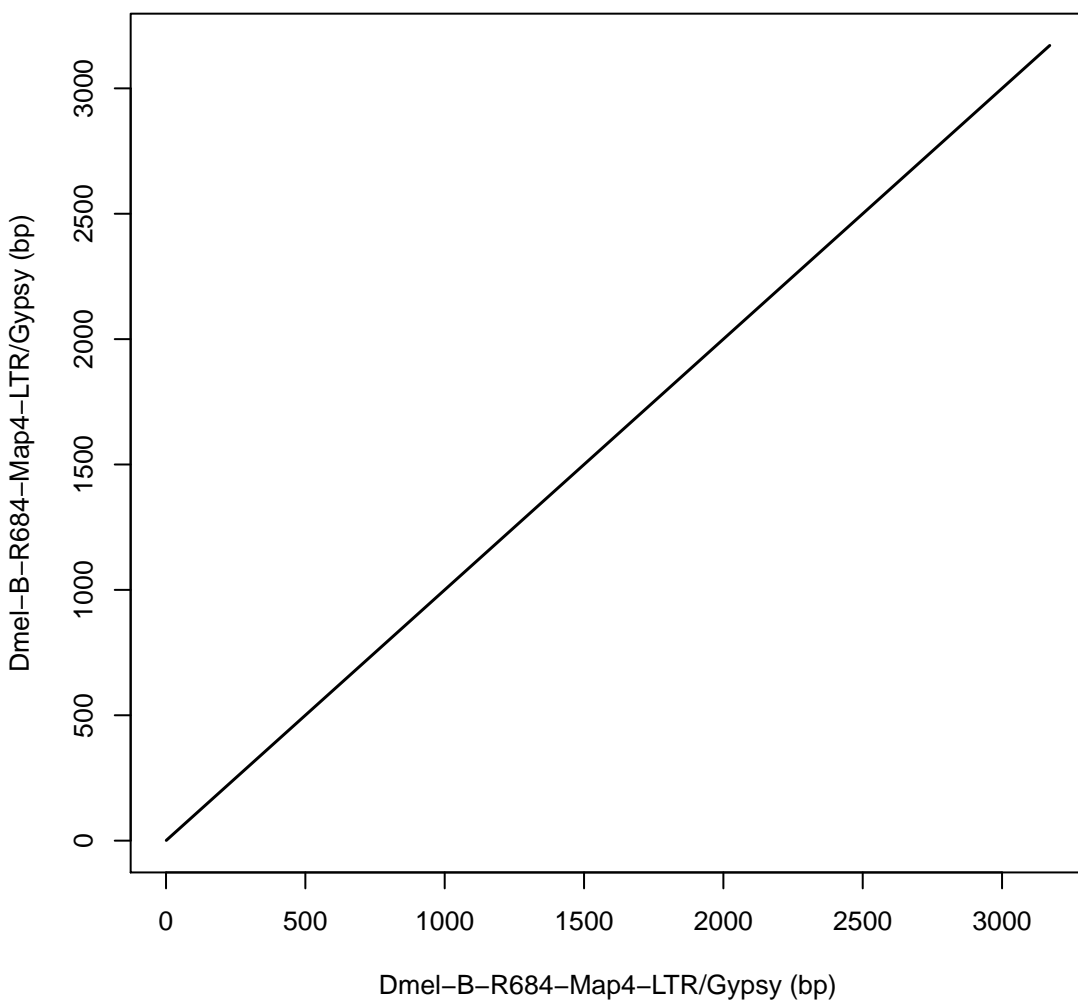
TE: Dmel-B-R684-Map4-LTR/Gypsy
consensus size: 3171bp; fragments: 195; full length: 0 (≥ 2853.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

