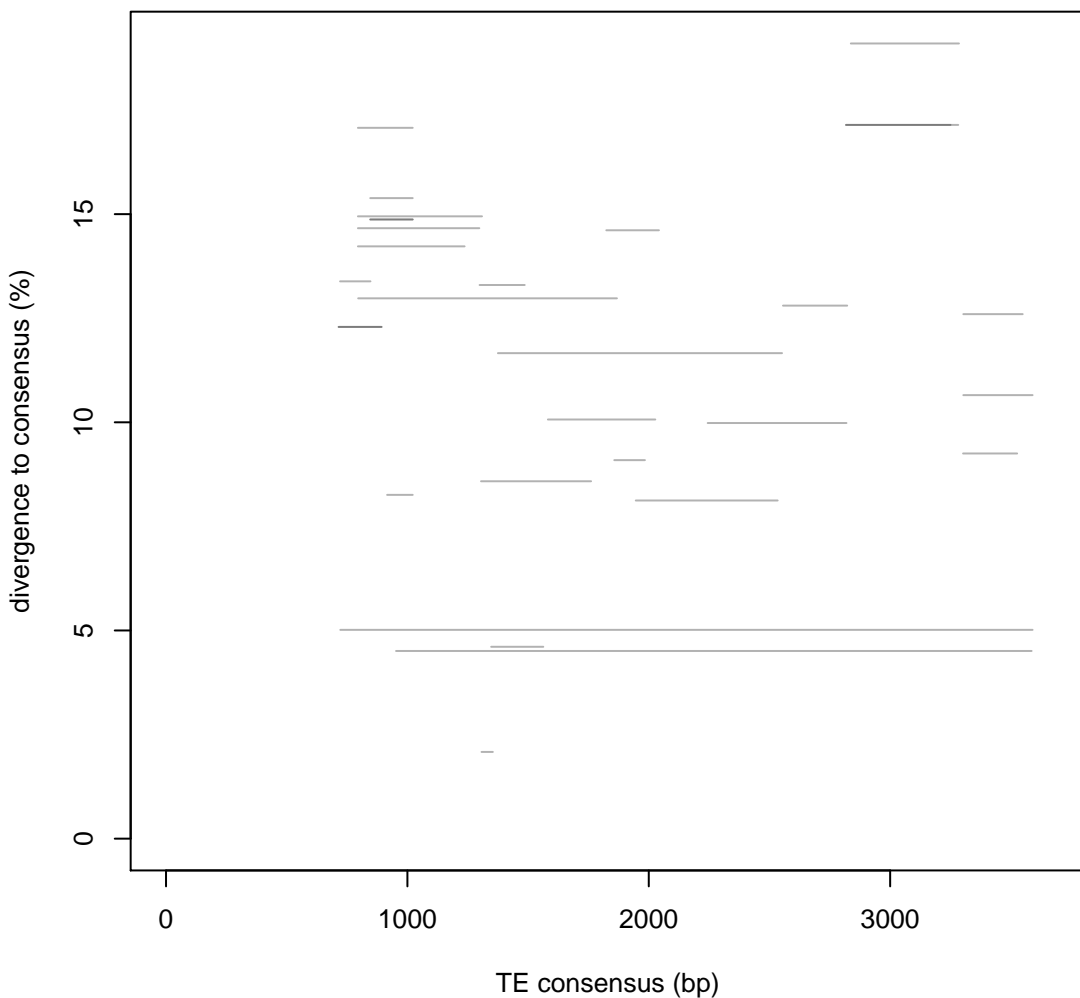
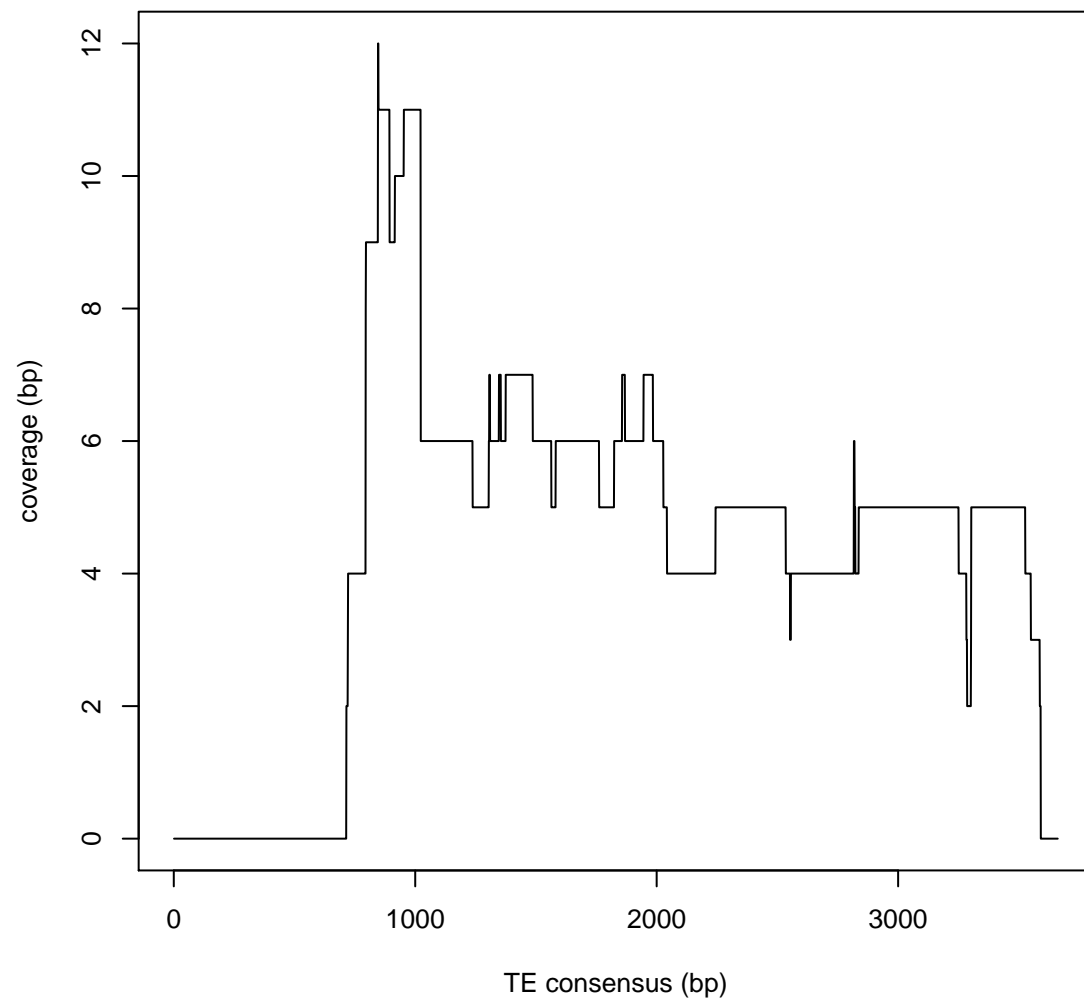


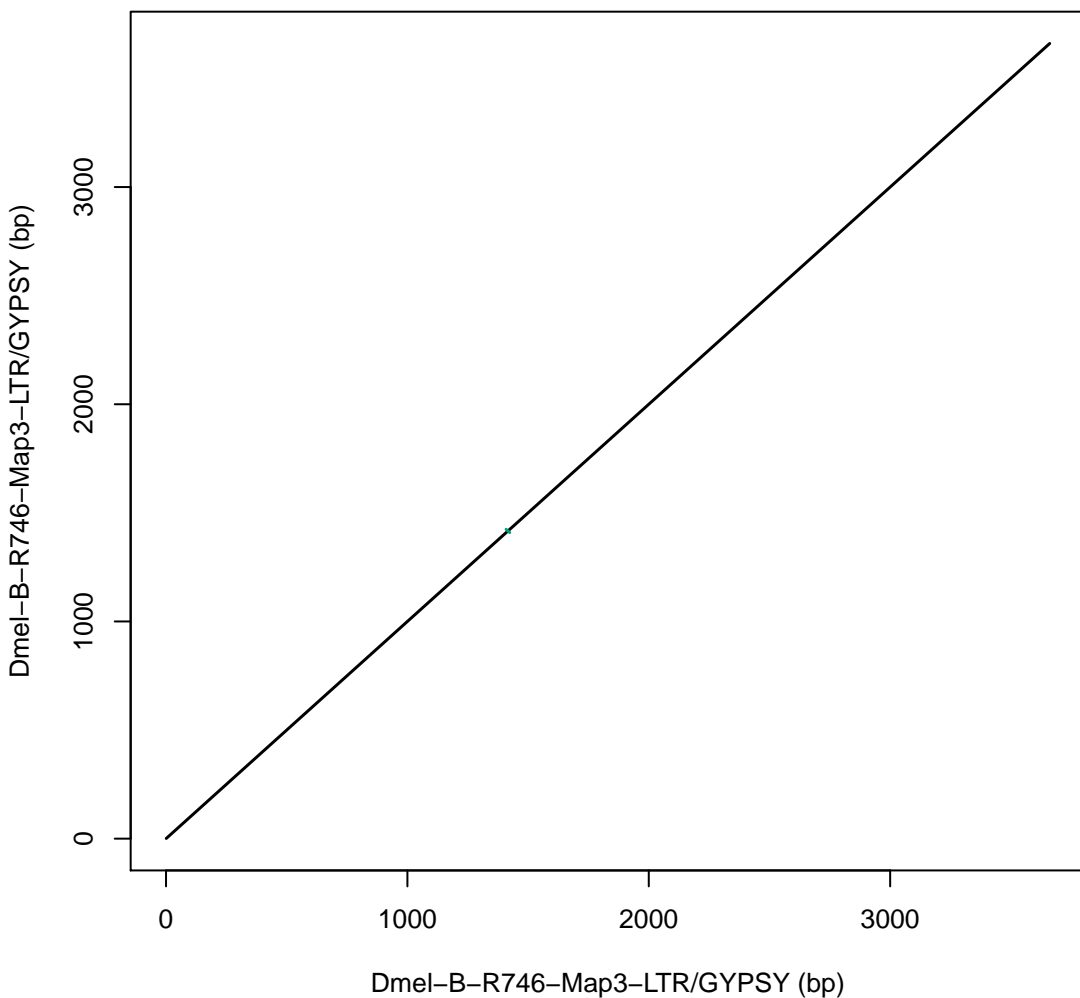
TE: Dmel-B-R746-Map3-LTR/GYPSY
consensus size: 3661bp; fragments: 31; full length: 0 (≥ 3294.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

