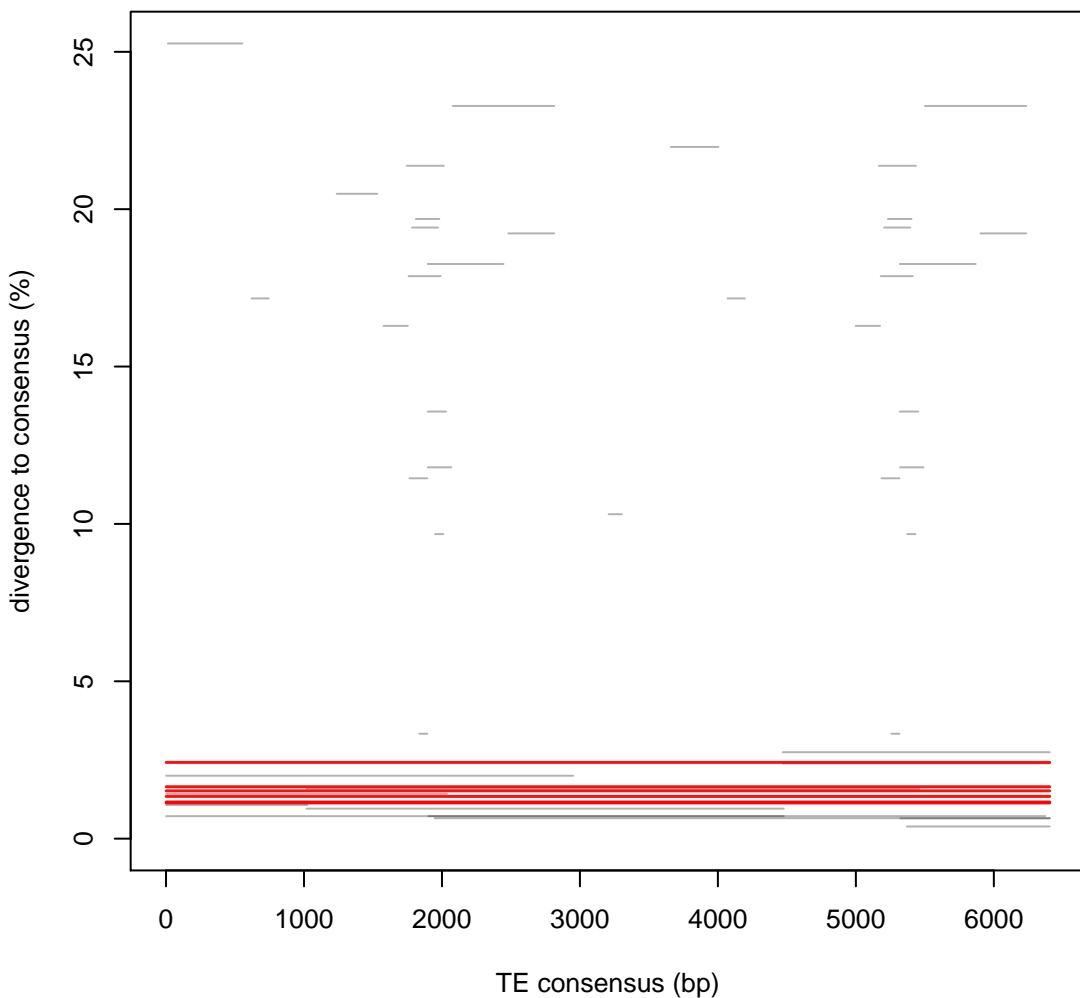
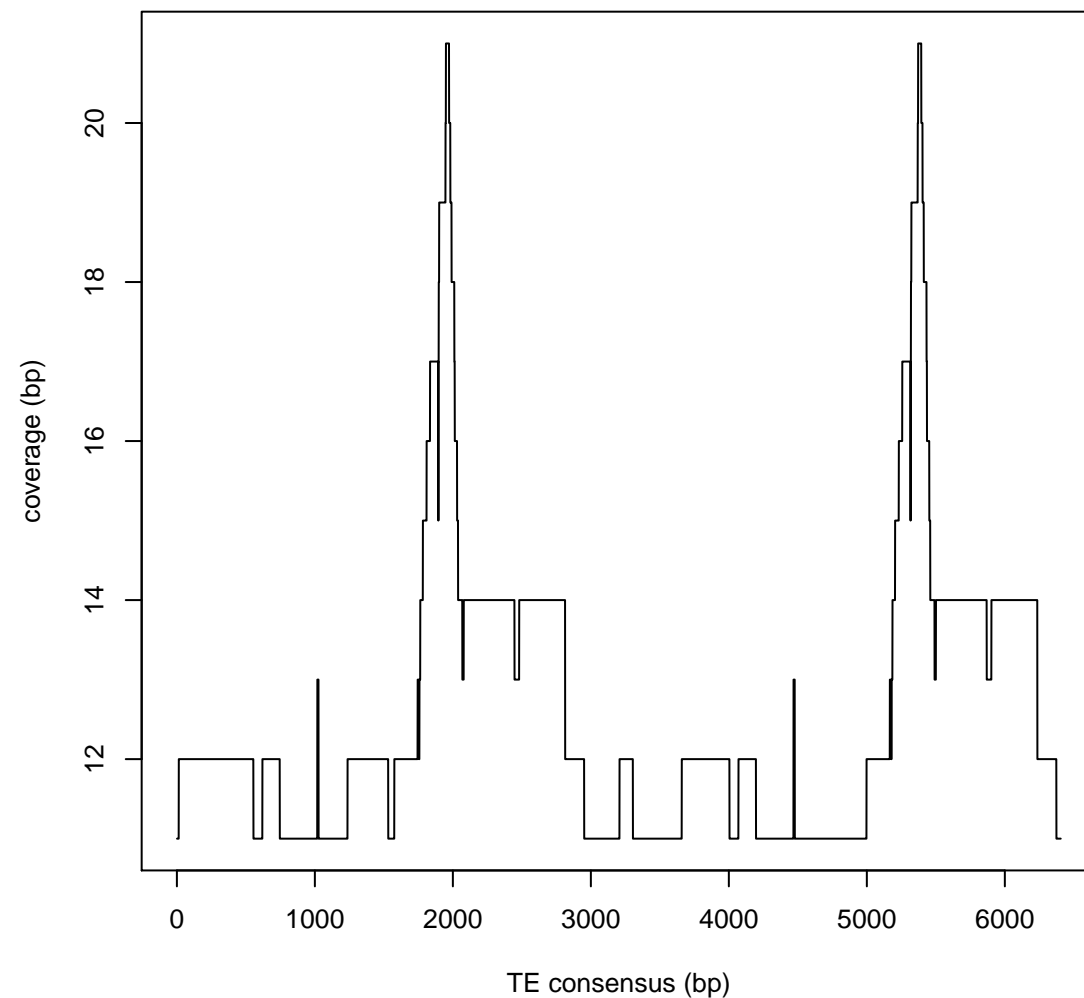


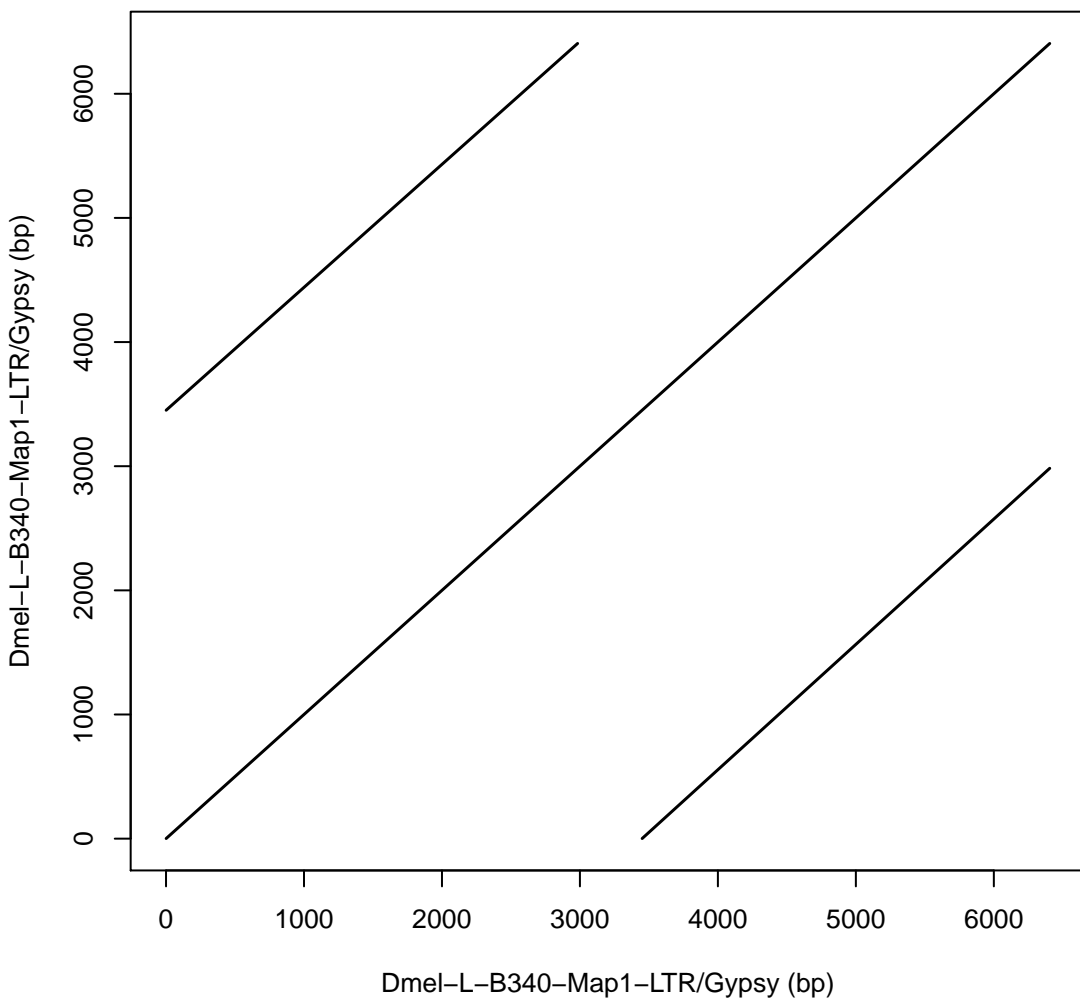
TE: Dmel-L-B340-Map1-LTR/Gypsy
consensus size: 6405bp; fragments: 51; full length: 6 (≥ 5764.5 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

