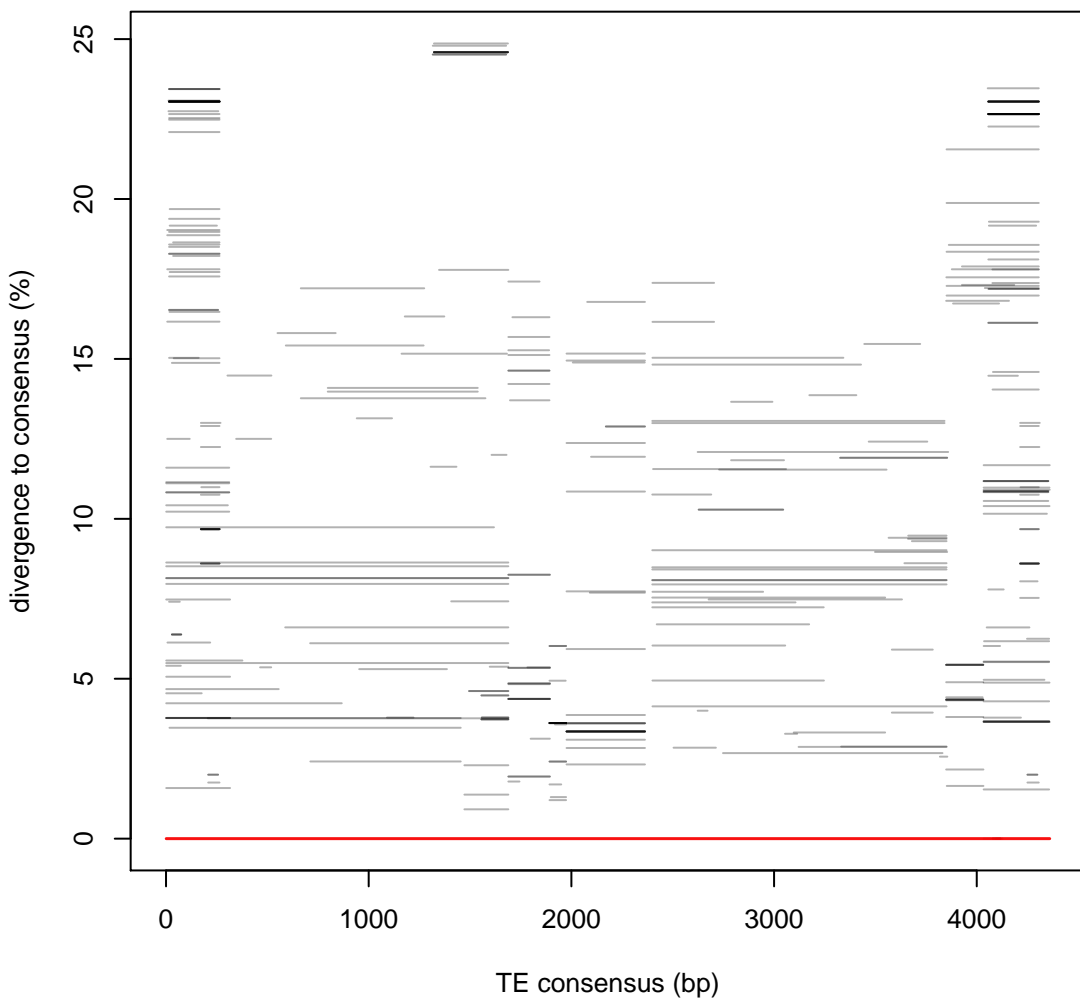
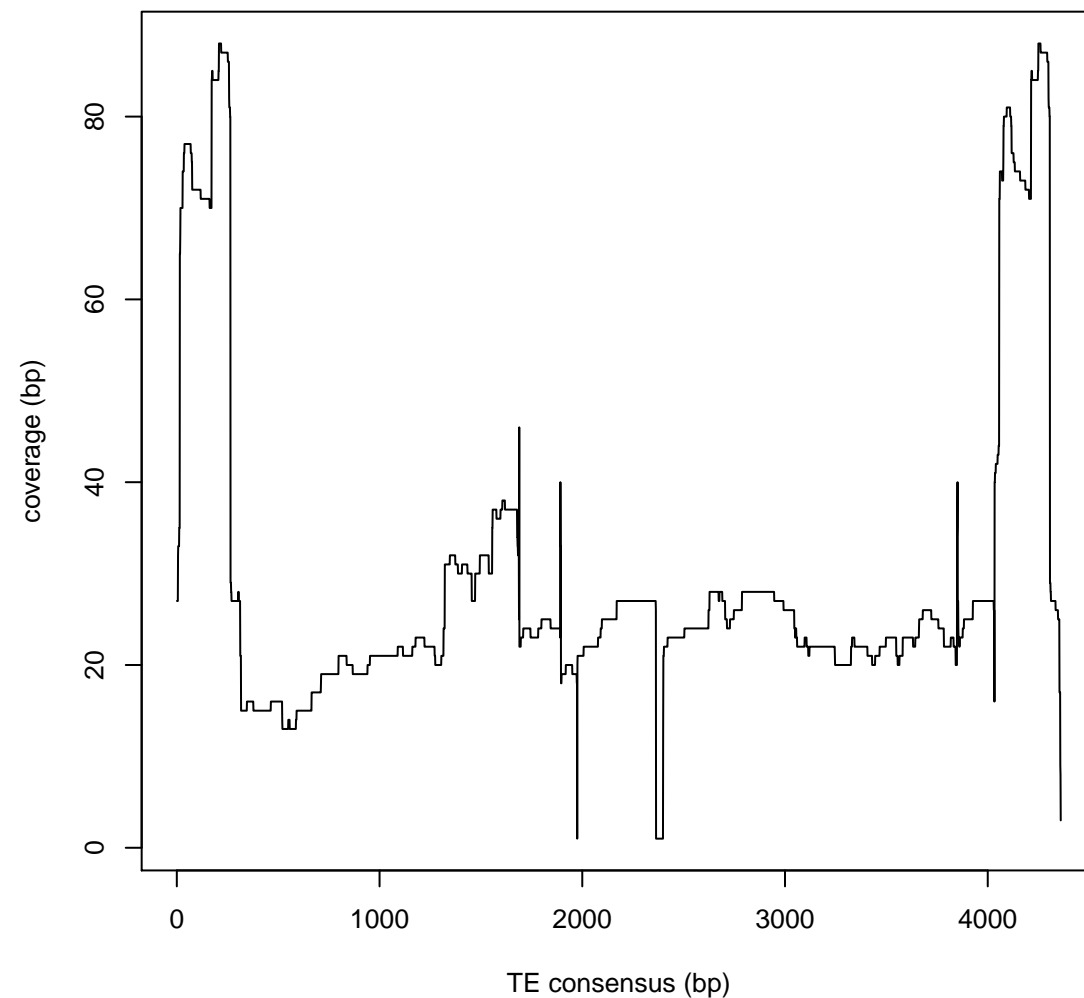


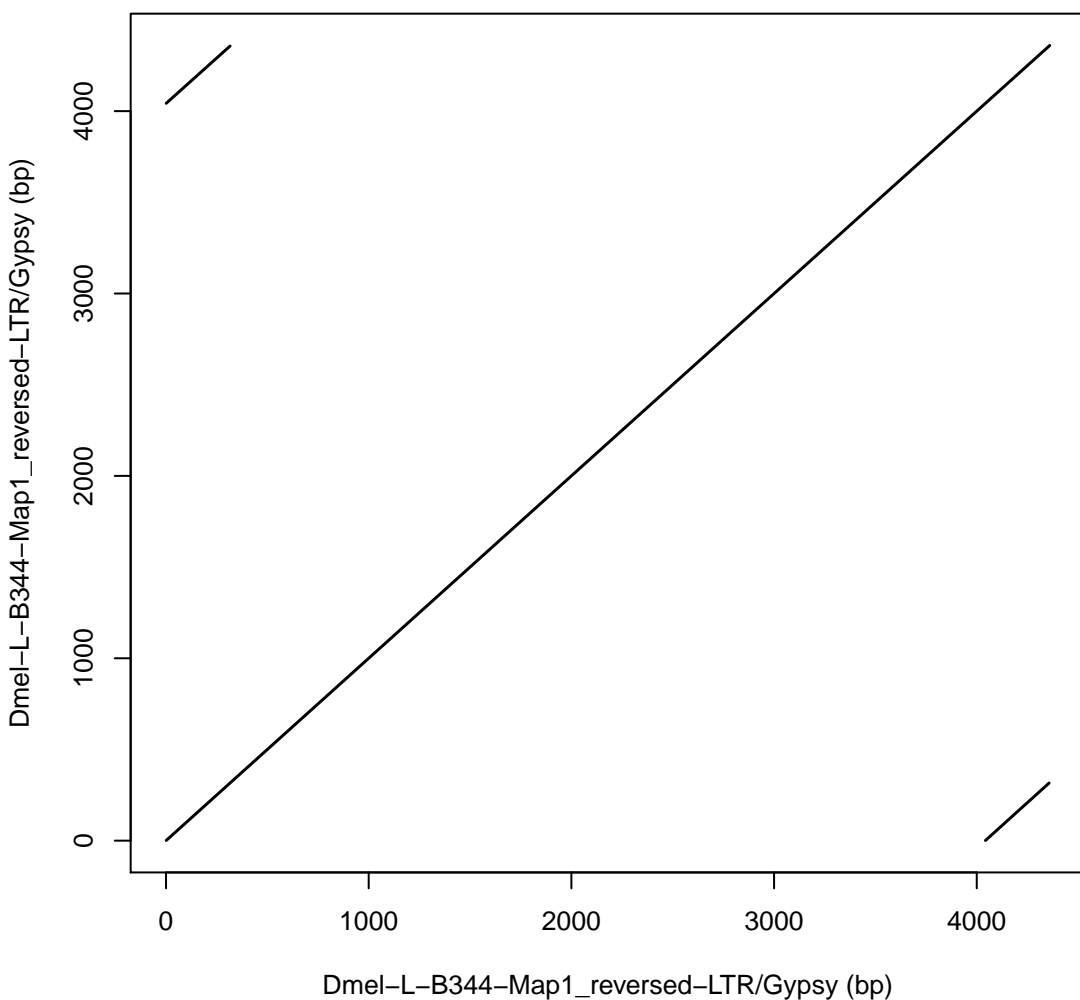
TE: Dmel-L-B344-Map1_reversed-LTR/Gypsy
consensus size: 4360bp; fragments: 373; full length: 1 (>=3924bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

