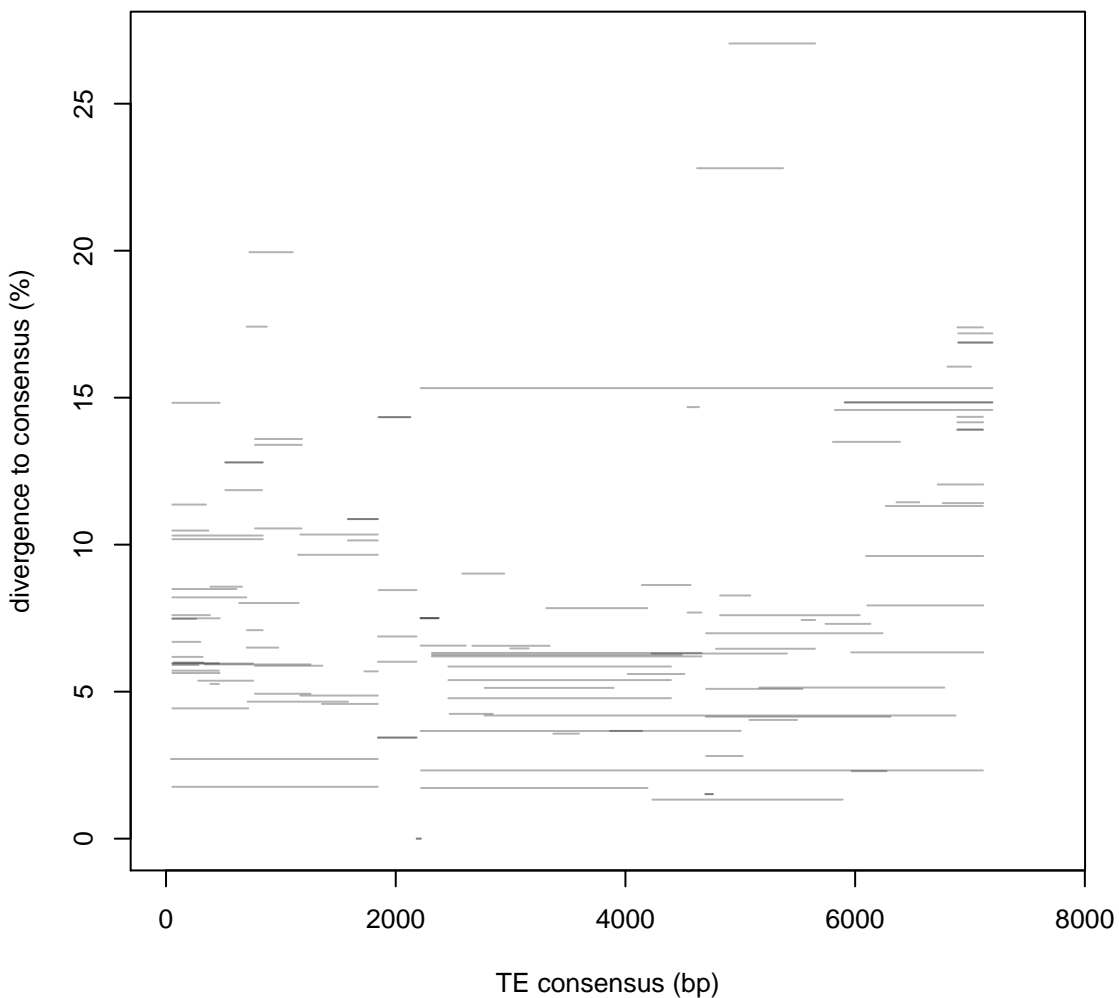
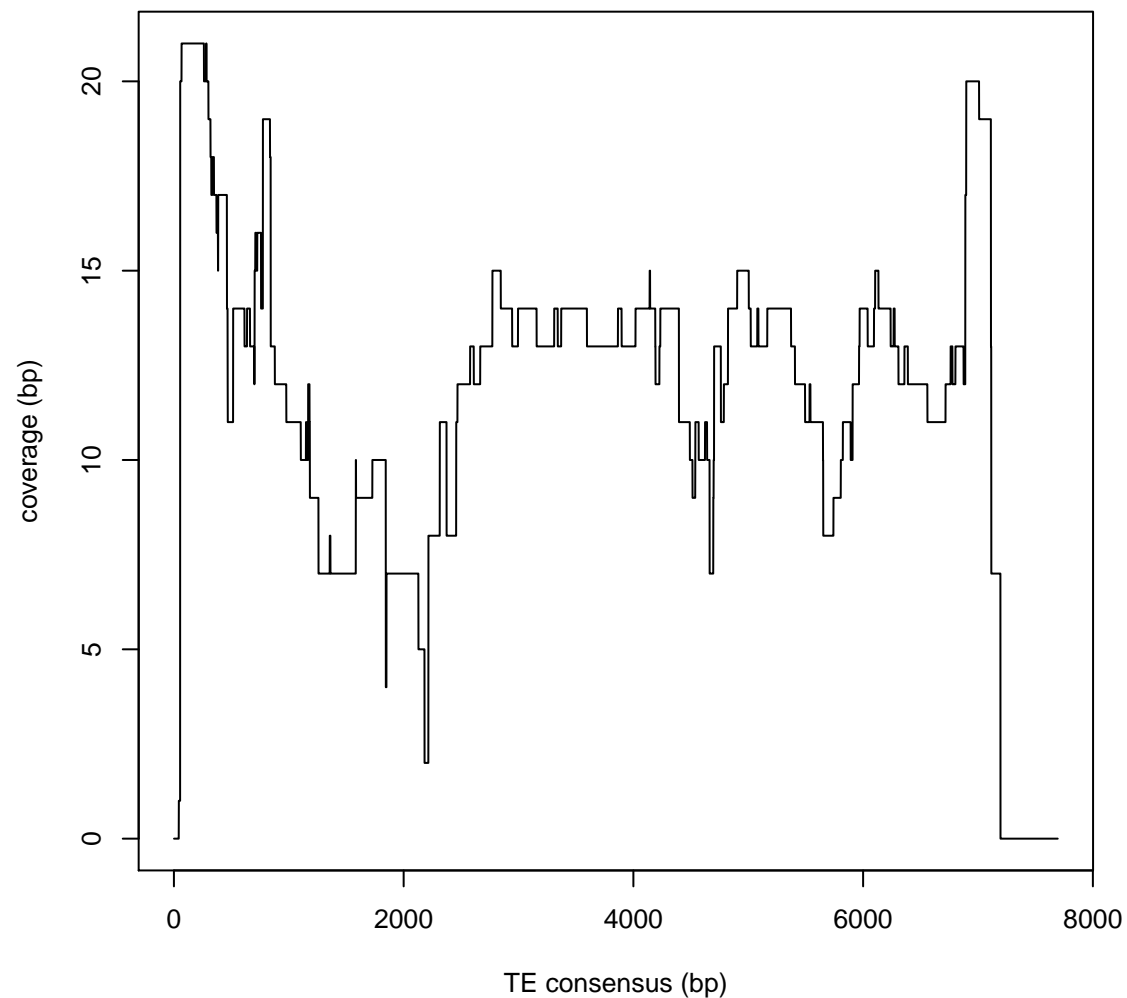


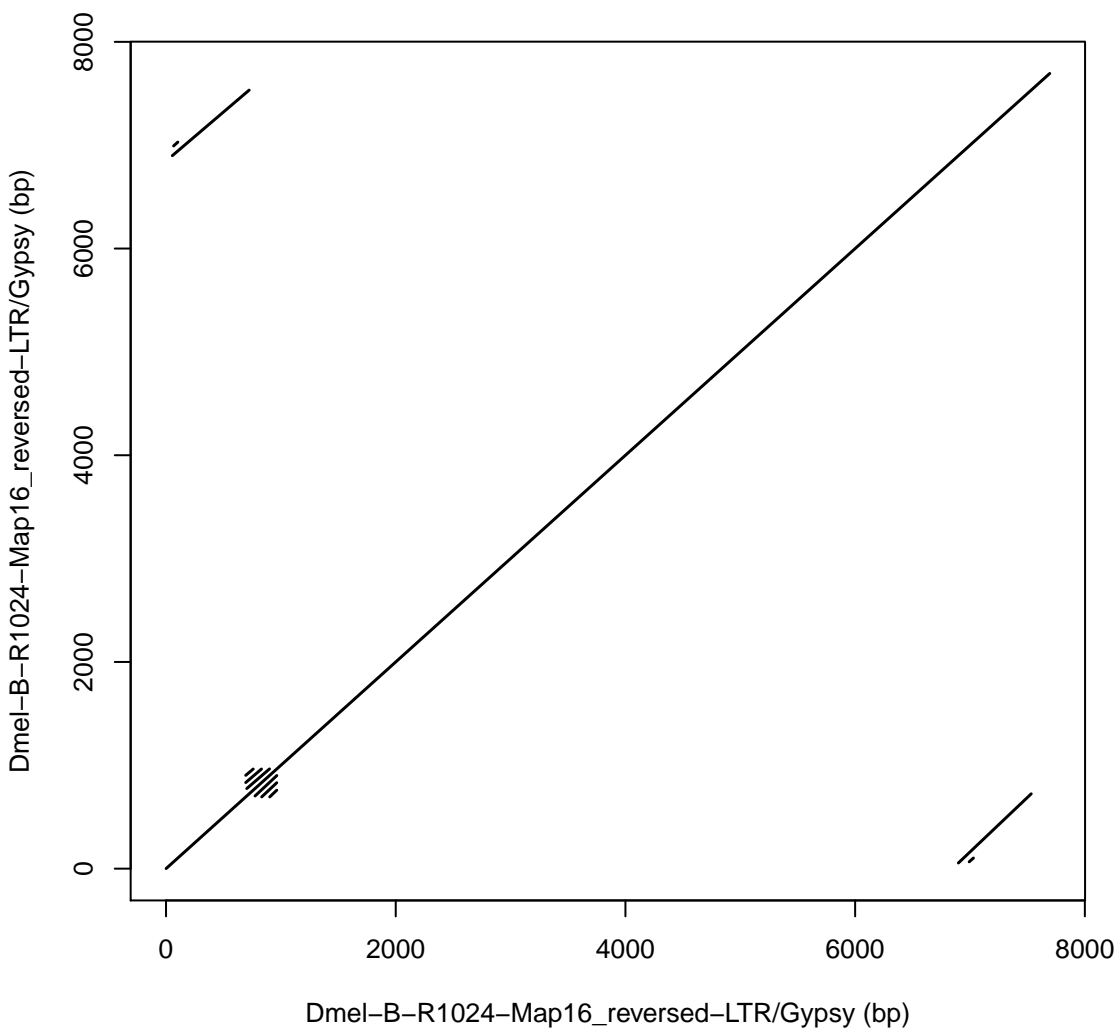
TE: Dmel-B-R1024-Map16_reversed-LTR/Gypsy
consensus size: 7694bp; fragments: 121; full length: 0 (>=6924.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

