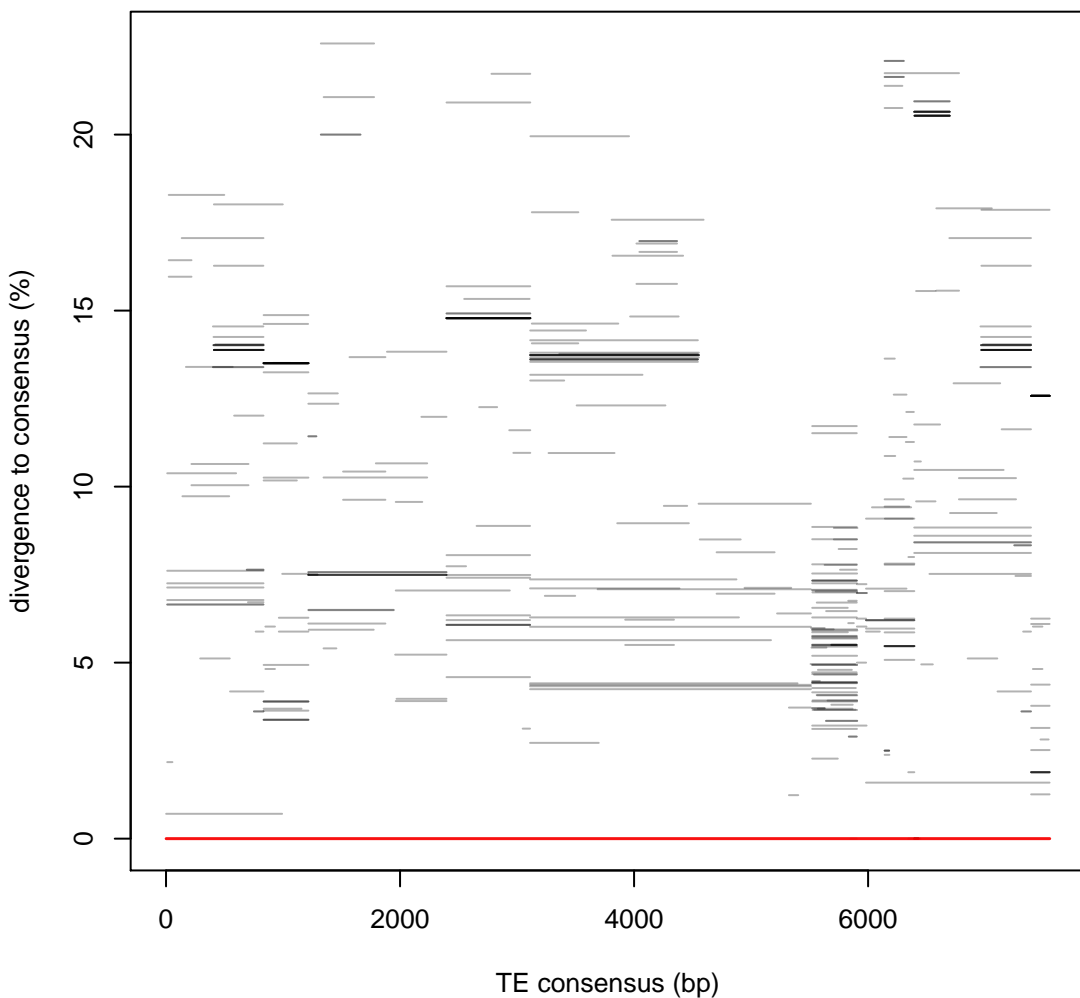
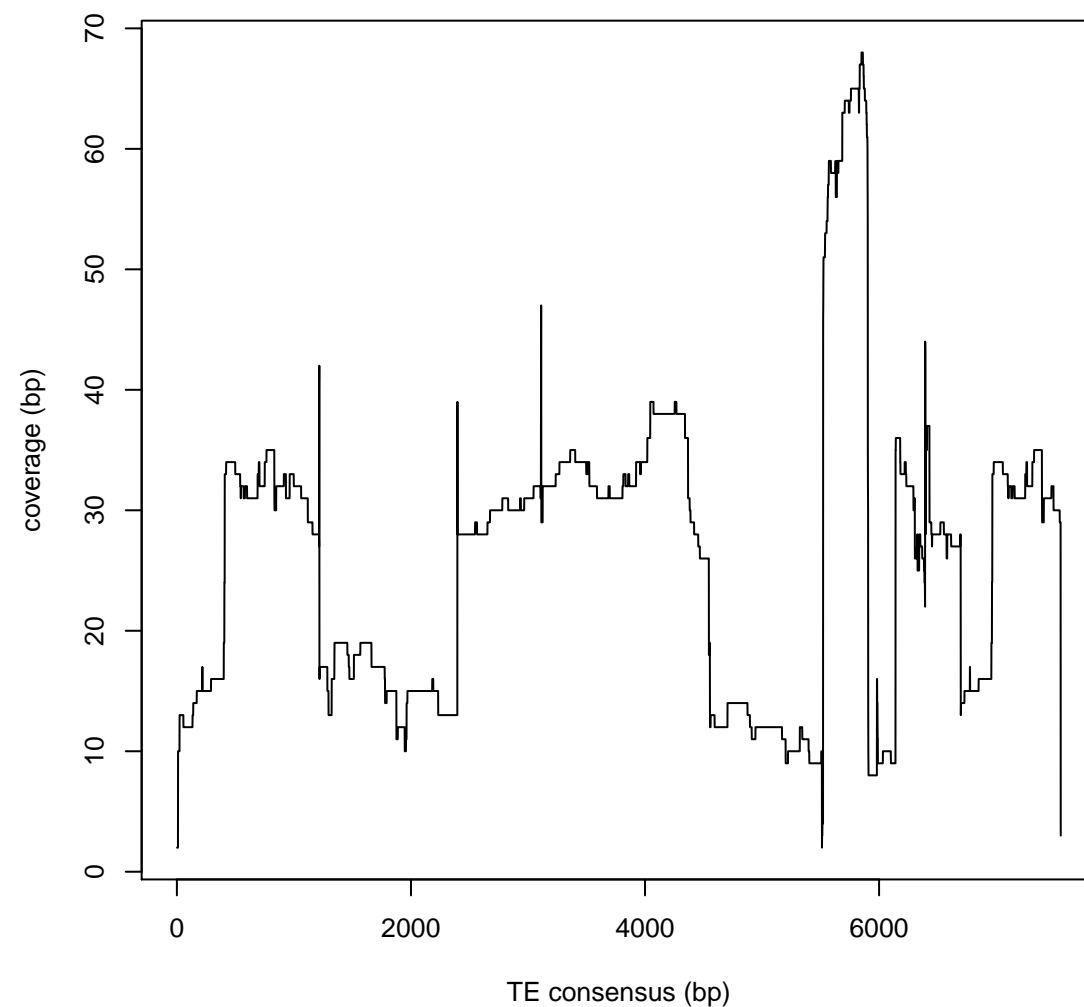


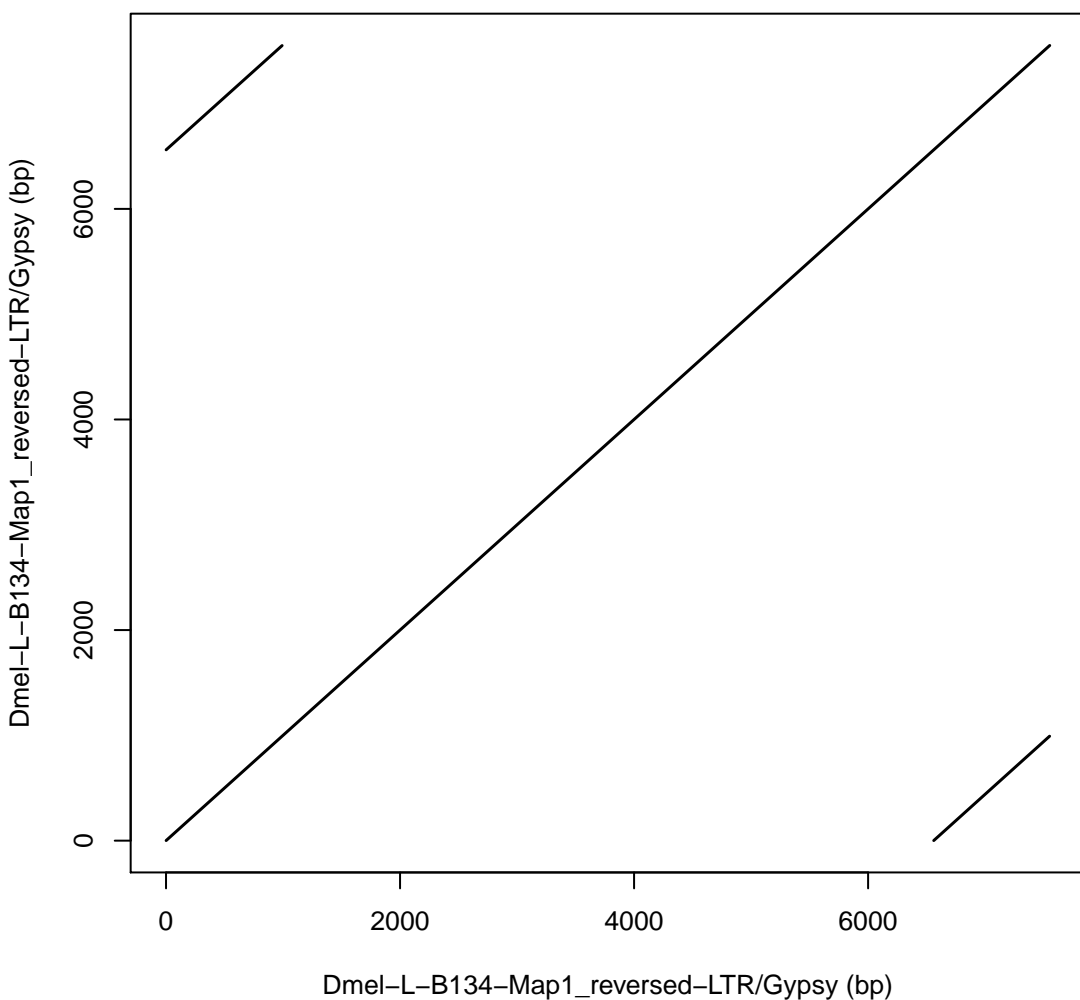
TE: Dmel-L-B134-Map1_reversed-LTR/Gypsy
consensus size: 7552bp; fragments: 424; full length: 1 (>=6796.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

