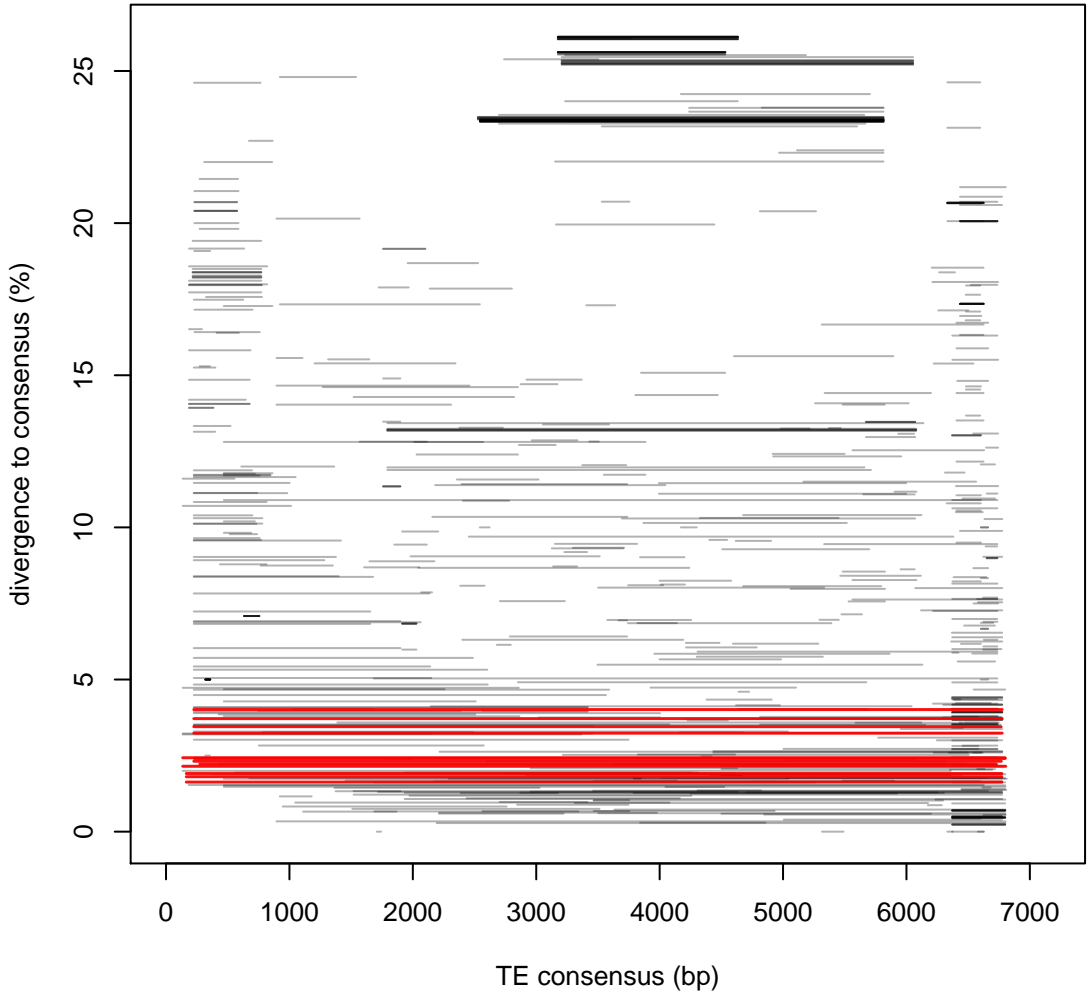
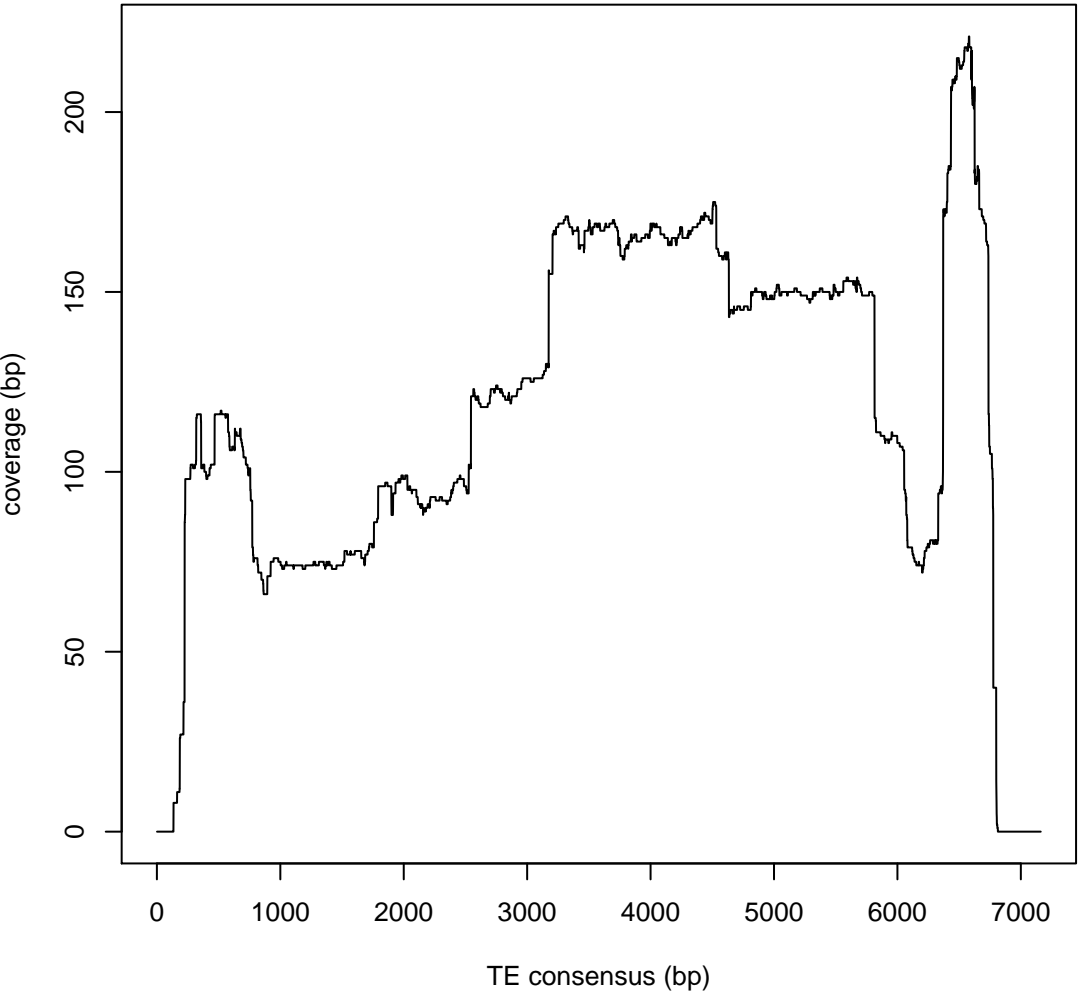


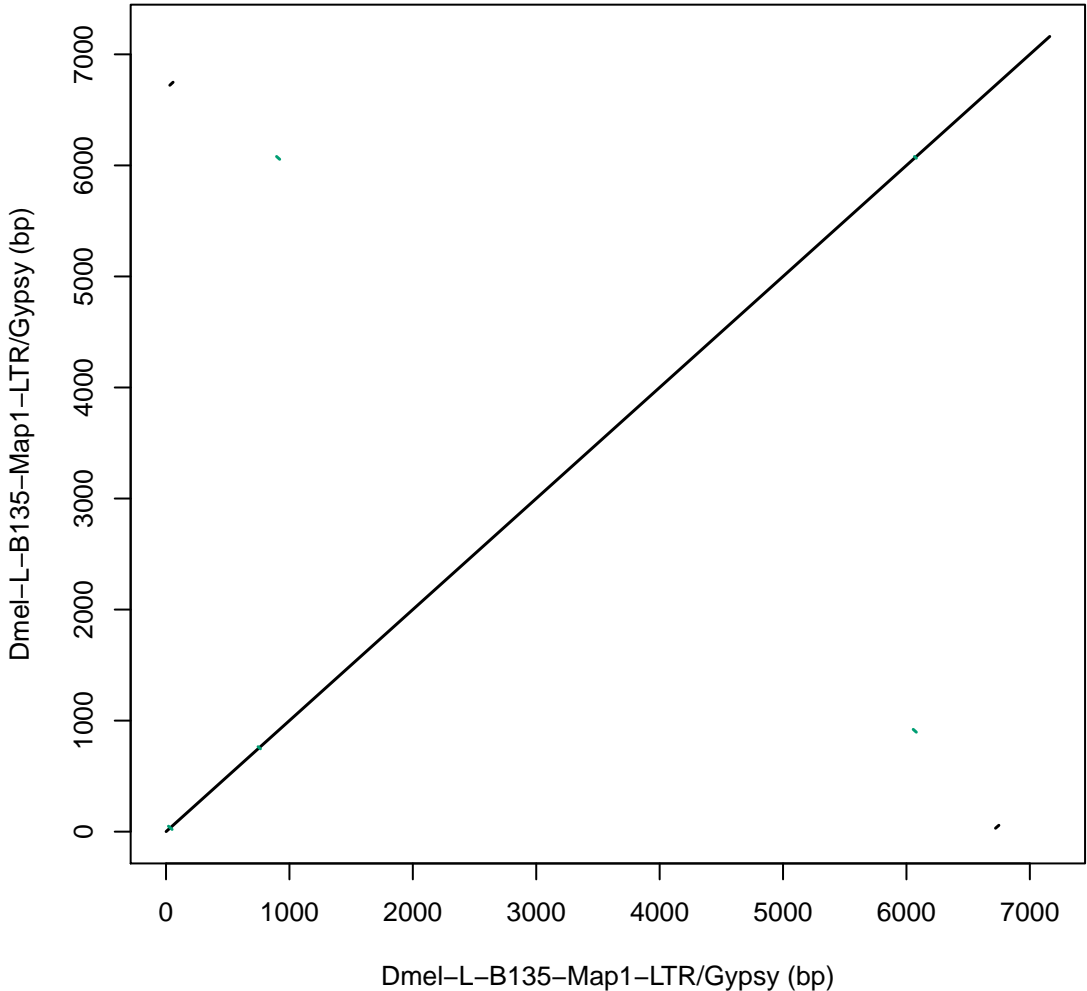
TE: Dmel-L-B135-Map1-LTR/Gypsy
consensus size: 7161bp; fragments: 658; full length: 12 (>=6444.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

