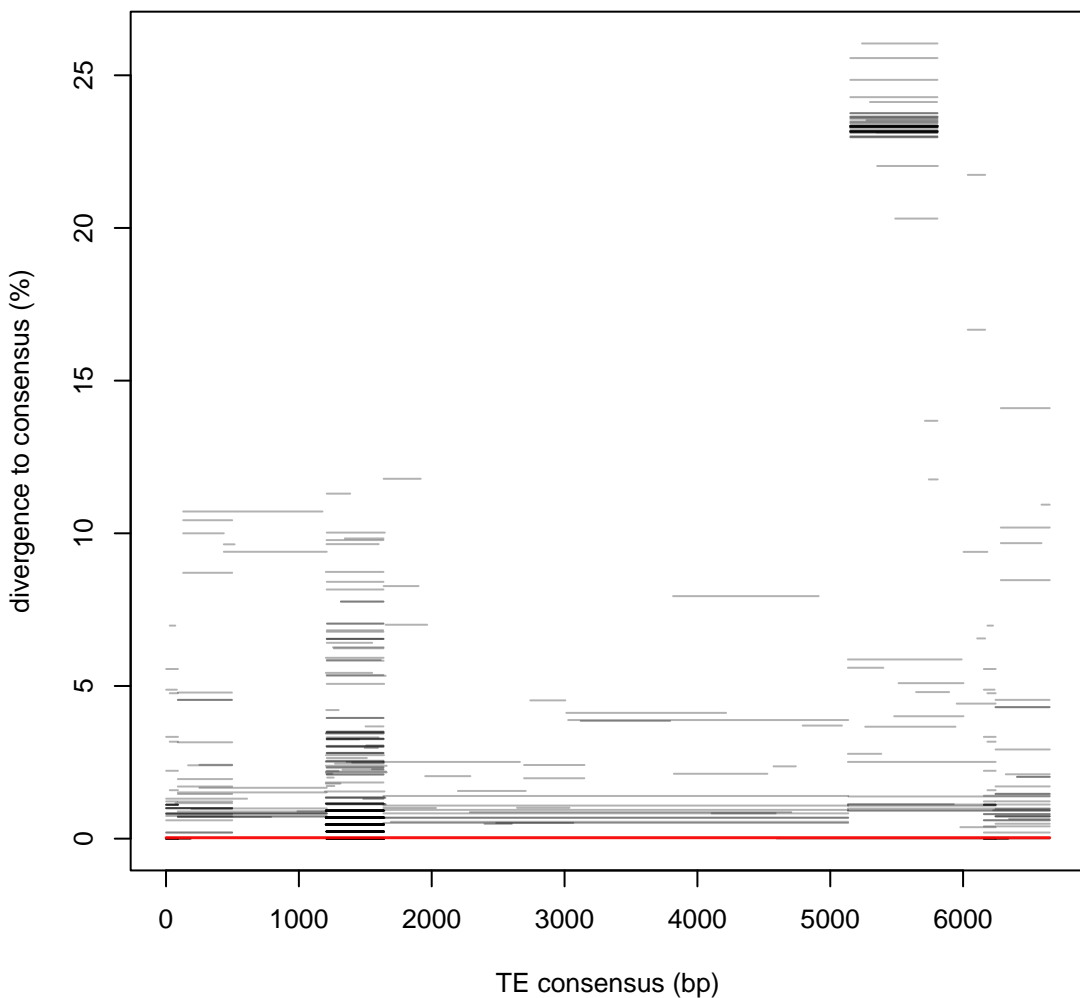
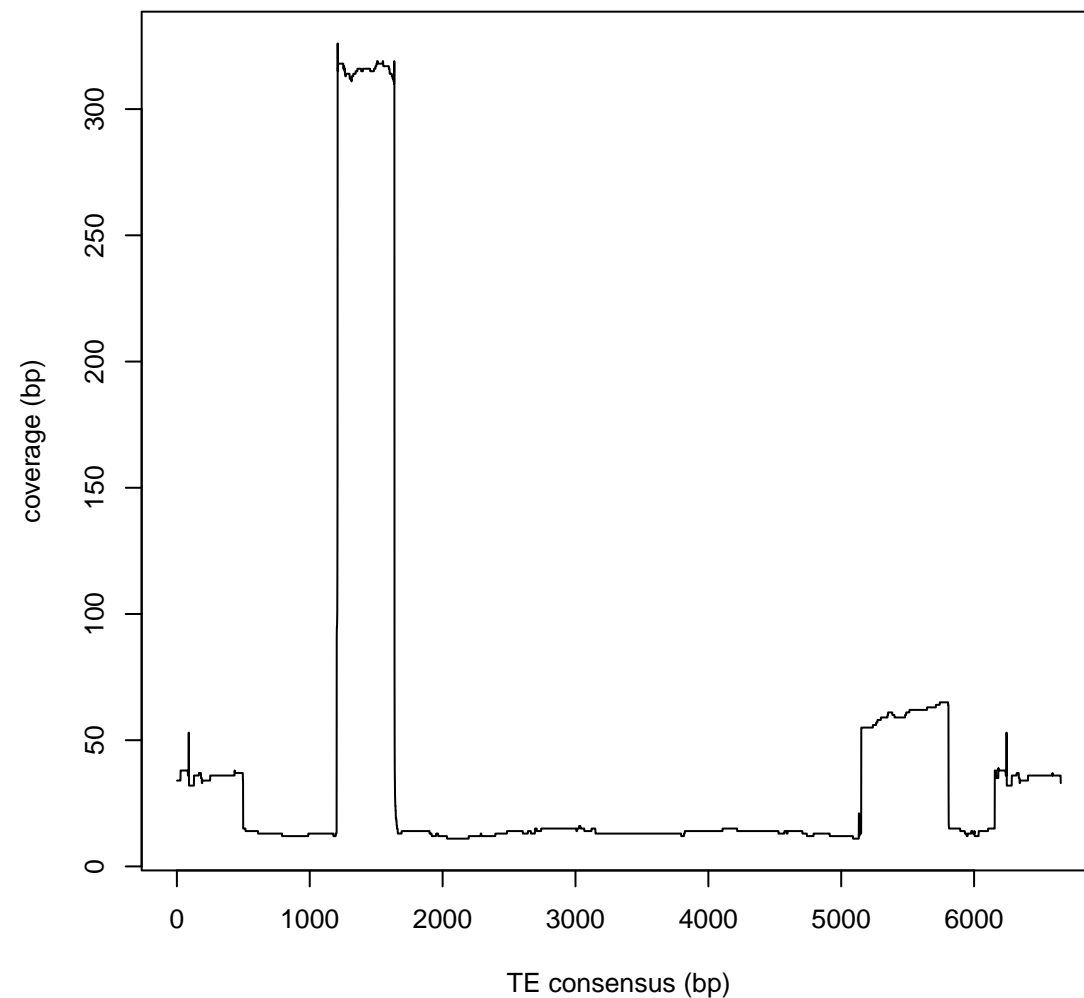


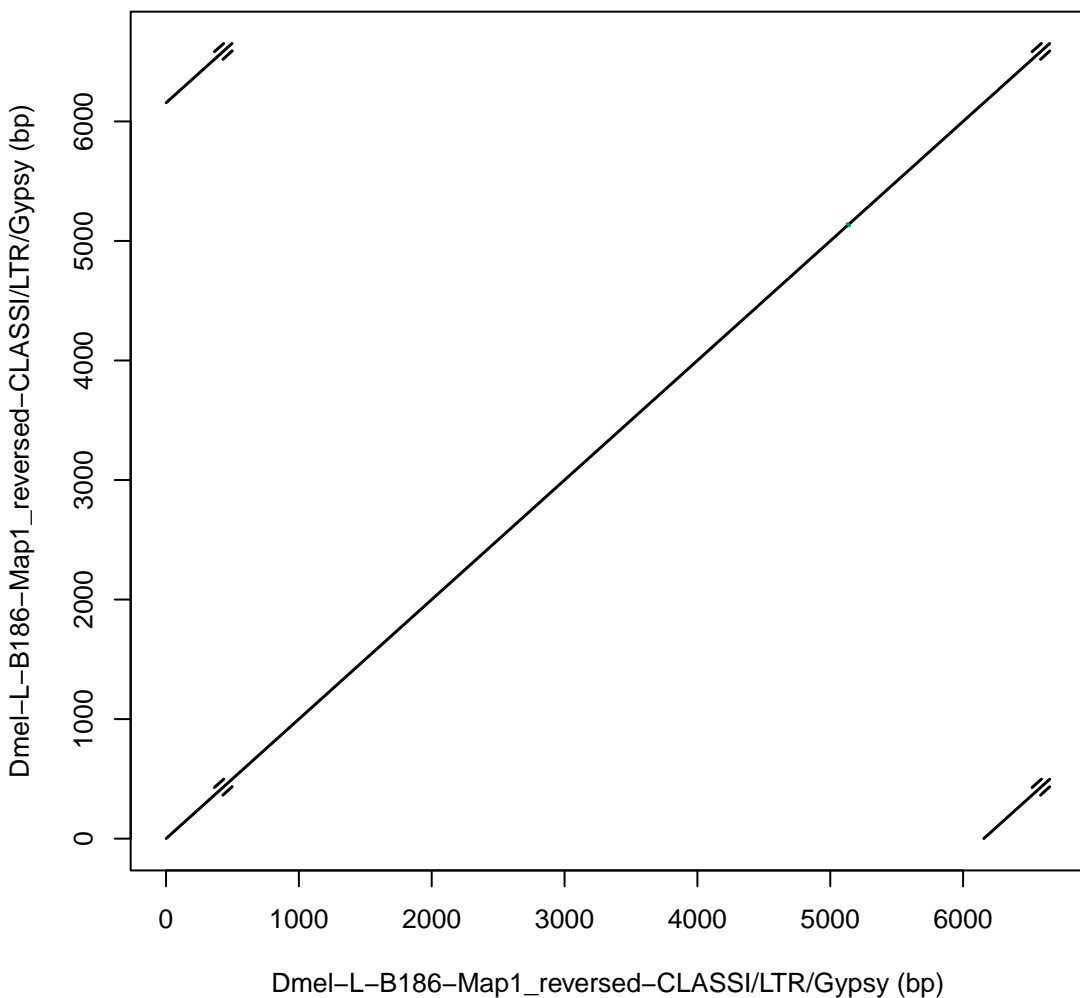
TE: Dmel-L-B186-Map1_reversed-CLASSI/LTR/Gypsy
consensus size: 6652bp; fragments: 554; full length: 1 (>=5986.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

