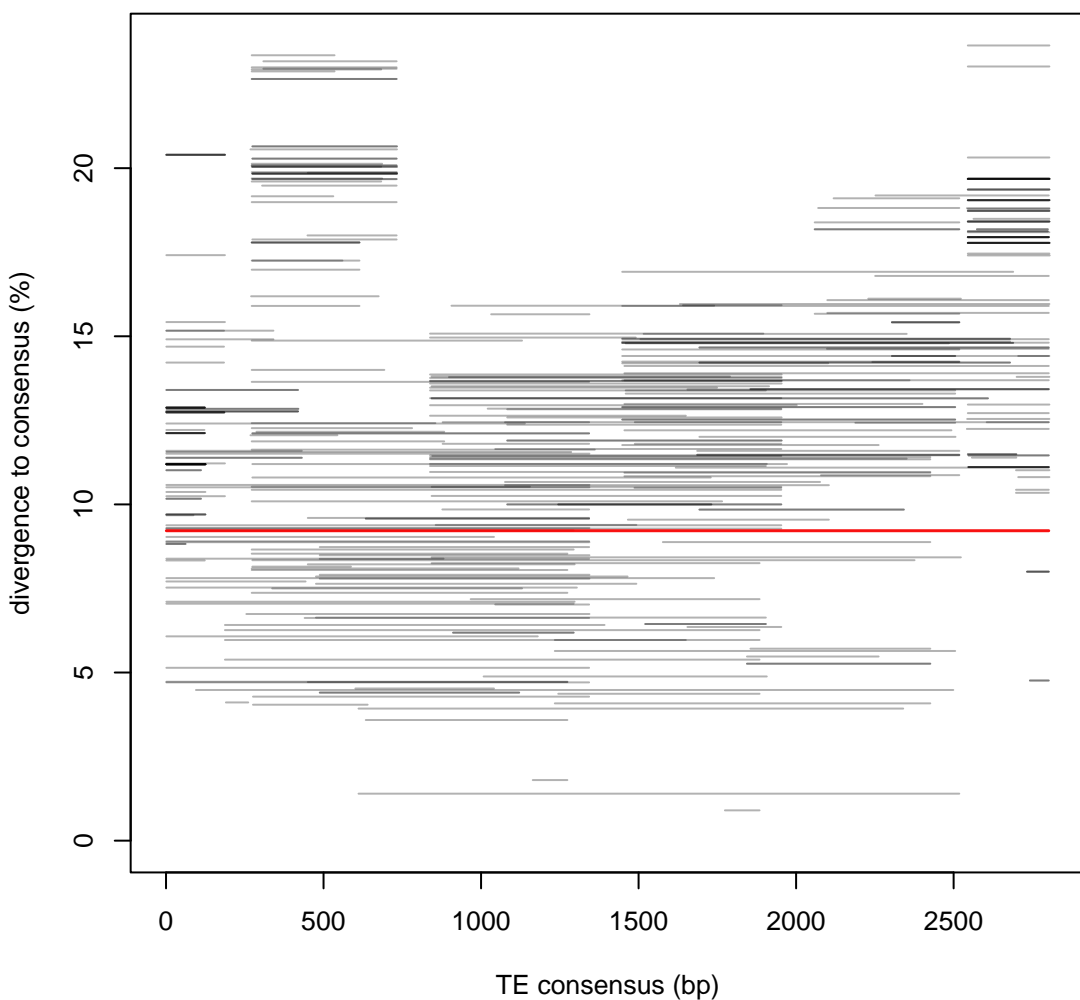
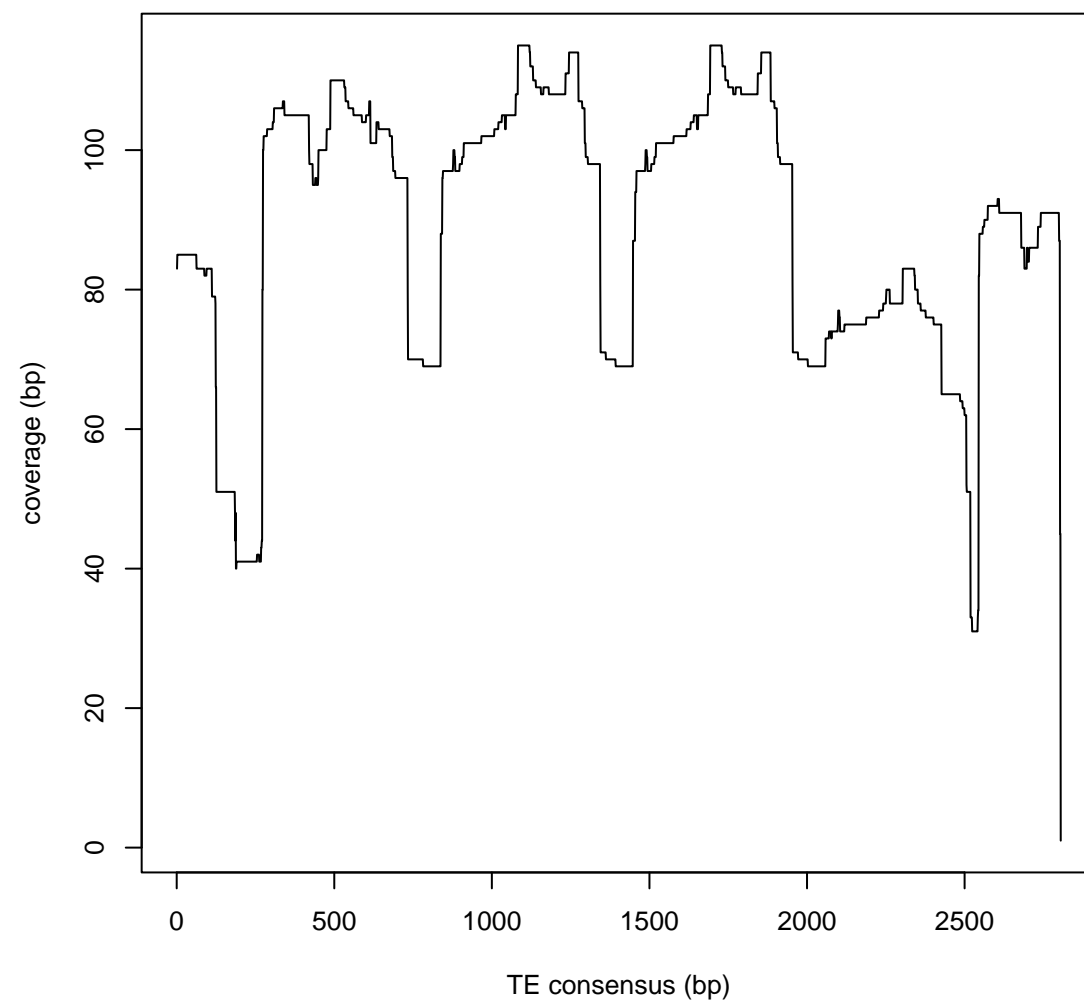


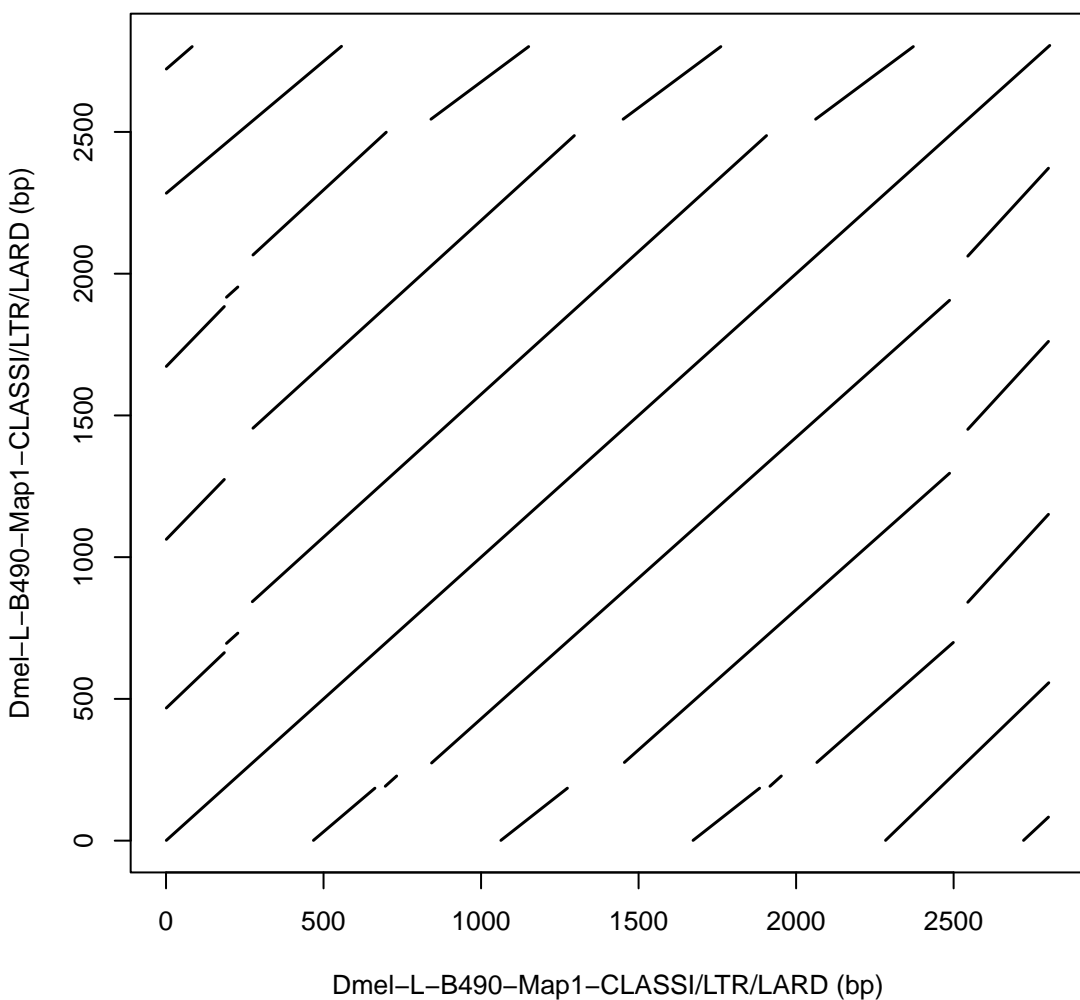
TE: Dmel-L-B490-Map1-CLASSI/LTR/LARD
consensus size: 2805bp; fragments: 391; full length: 1 (≥ 2524.5 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

