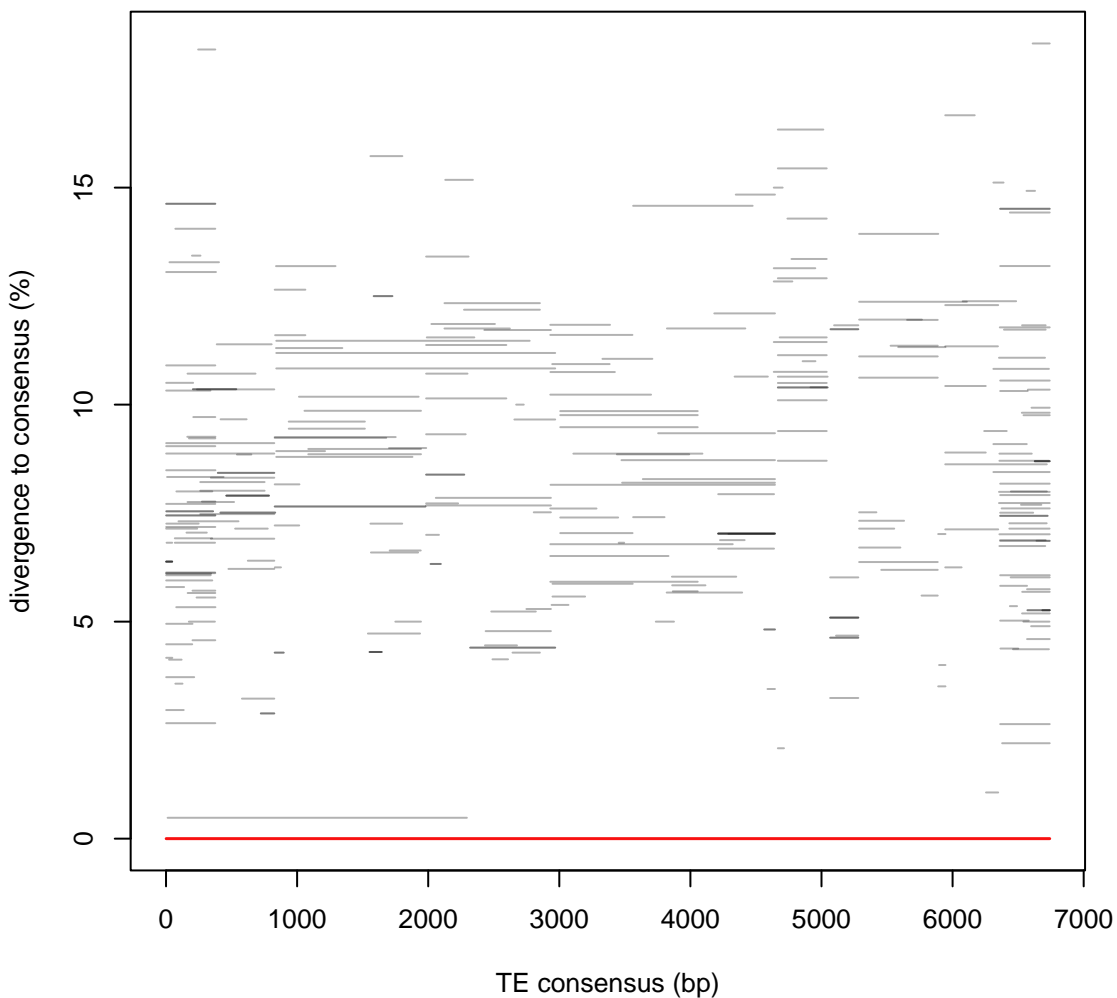
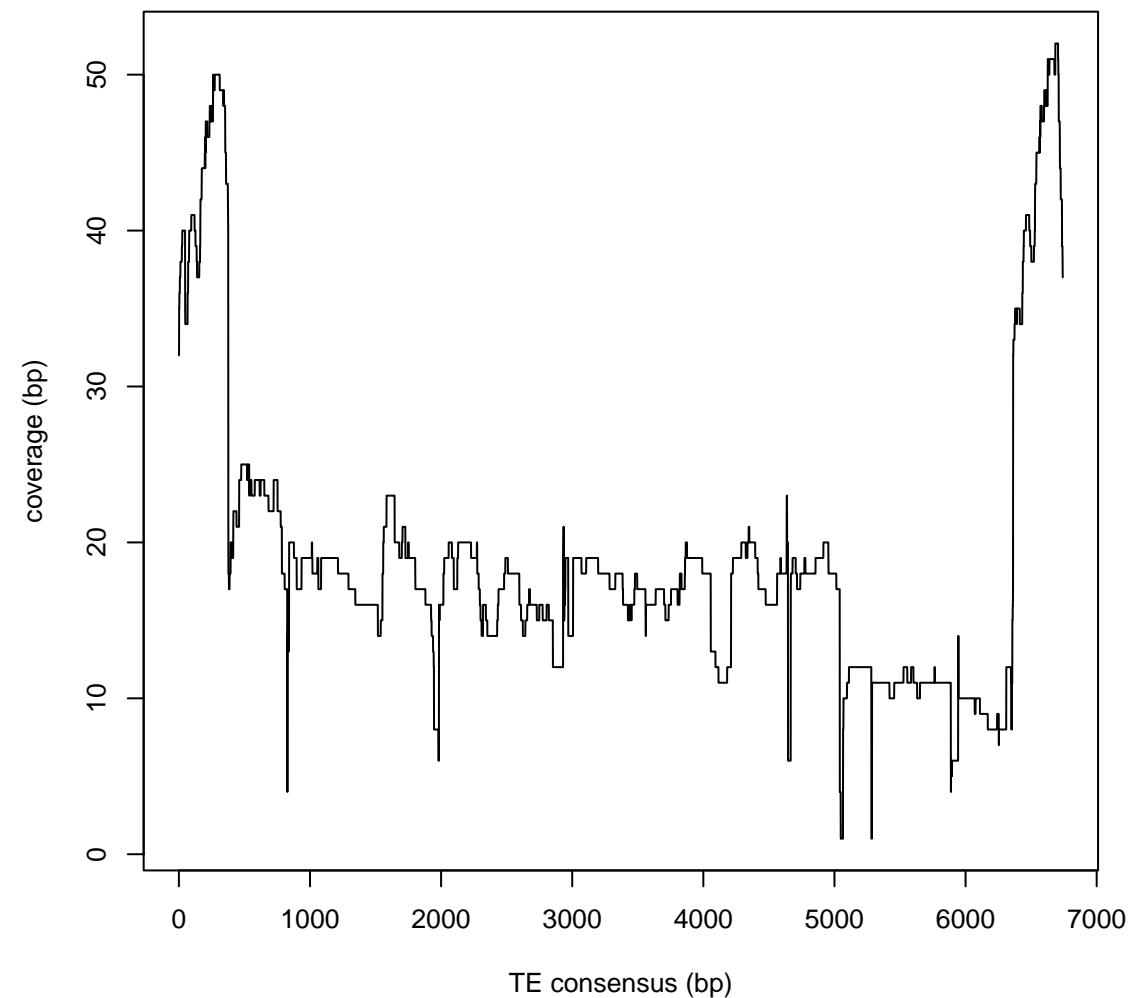


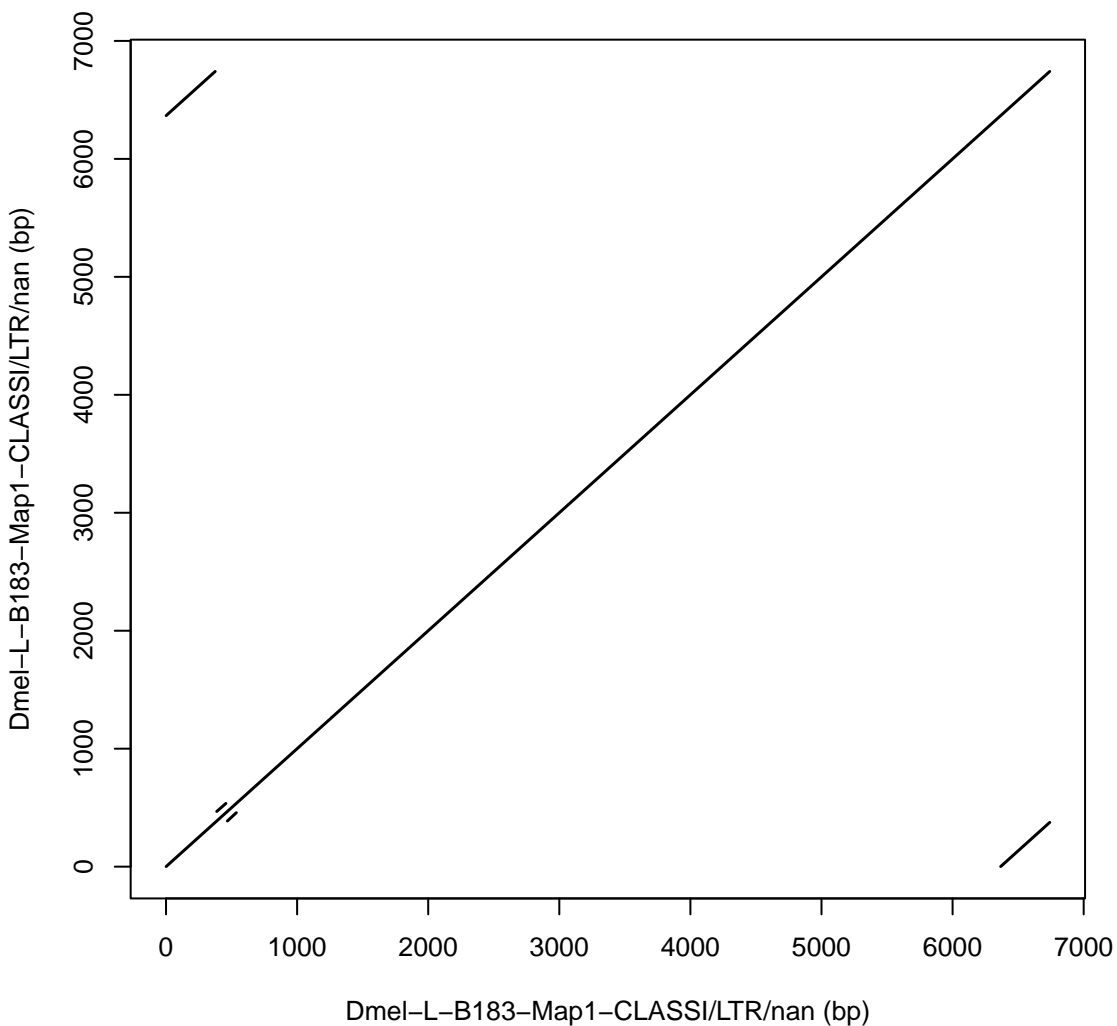
TE: Dmel-L-B183-Map1-CLASSI/LTR/nan  
consensus size: 6741bp; fragments: 323; full length: 1 (>=6066.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

