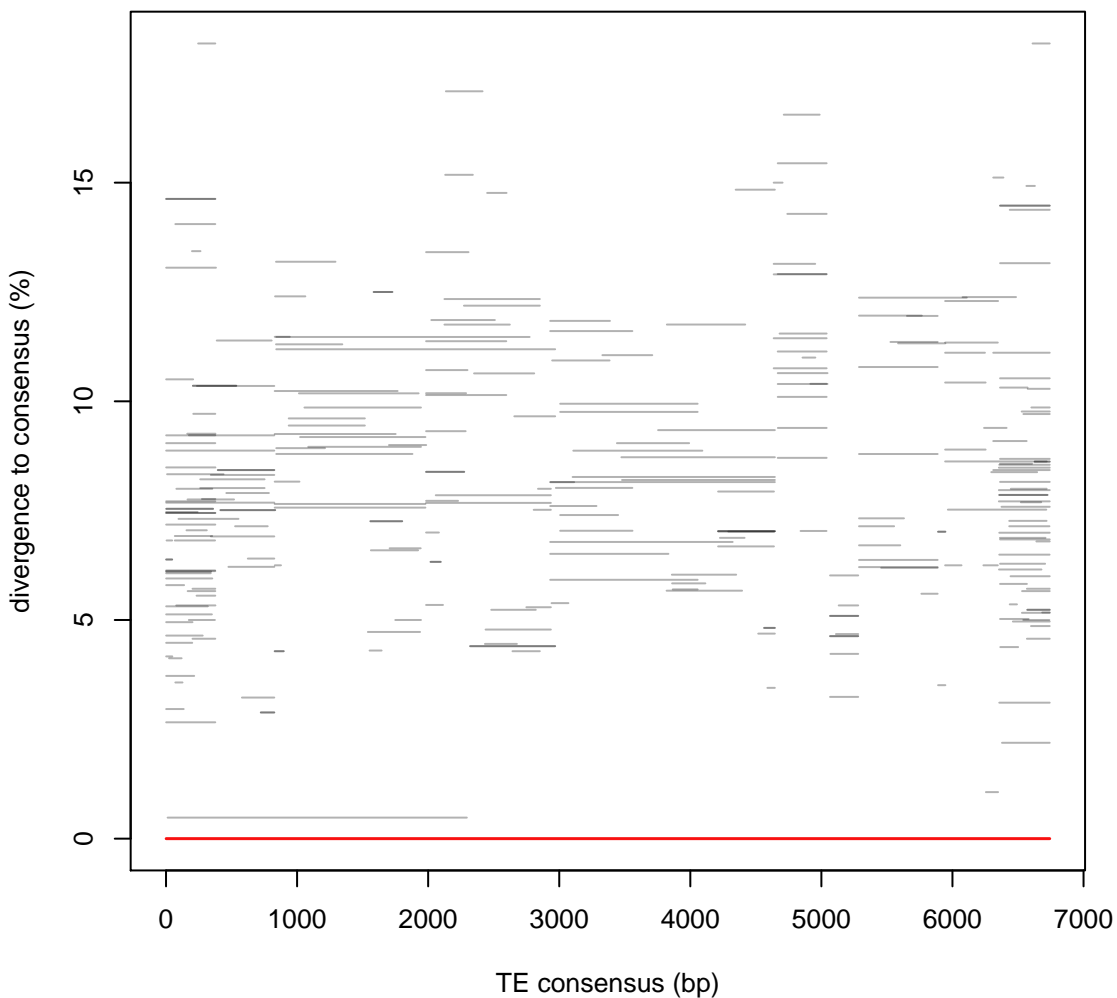
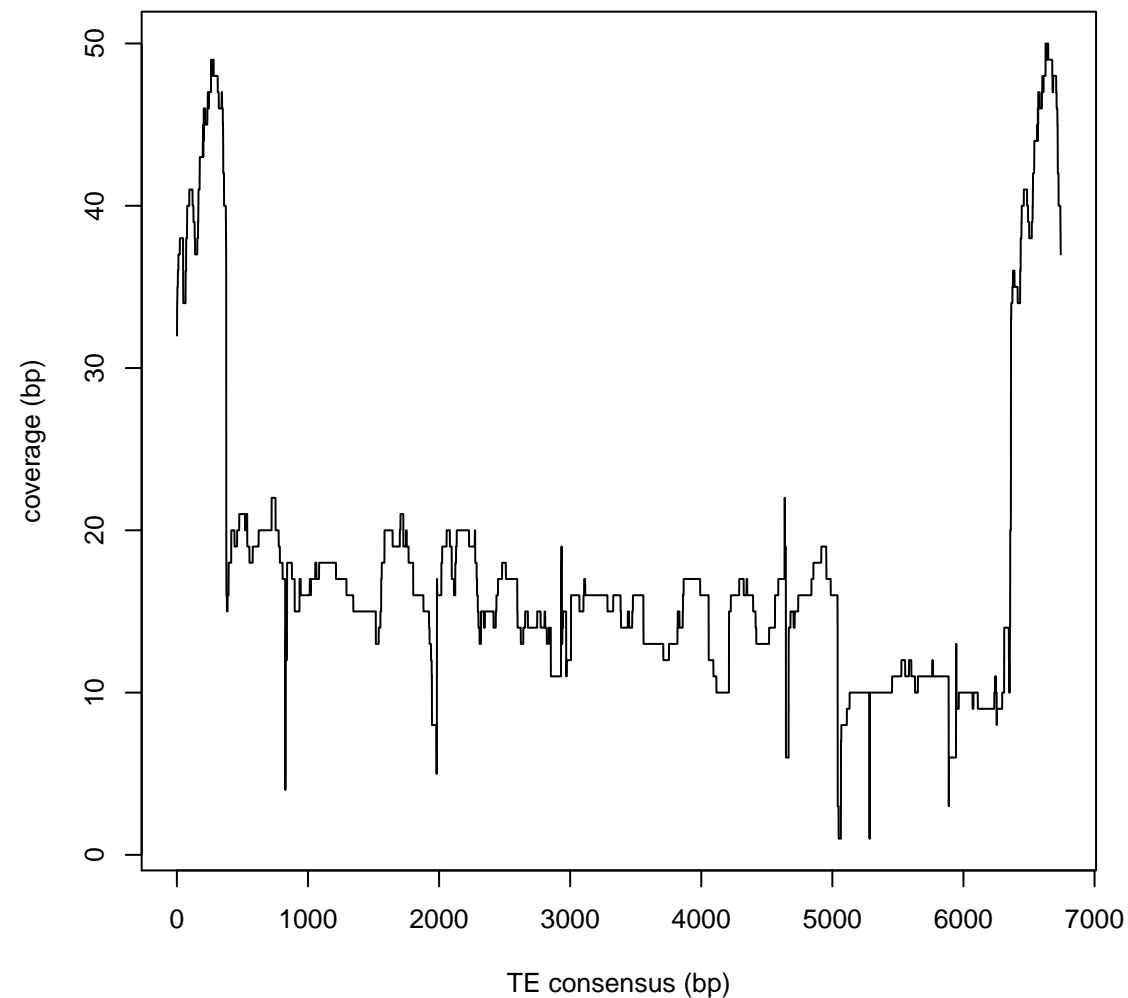


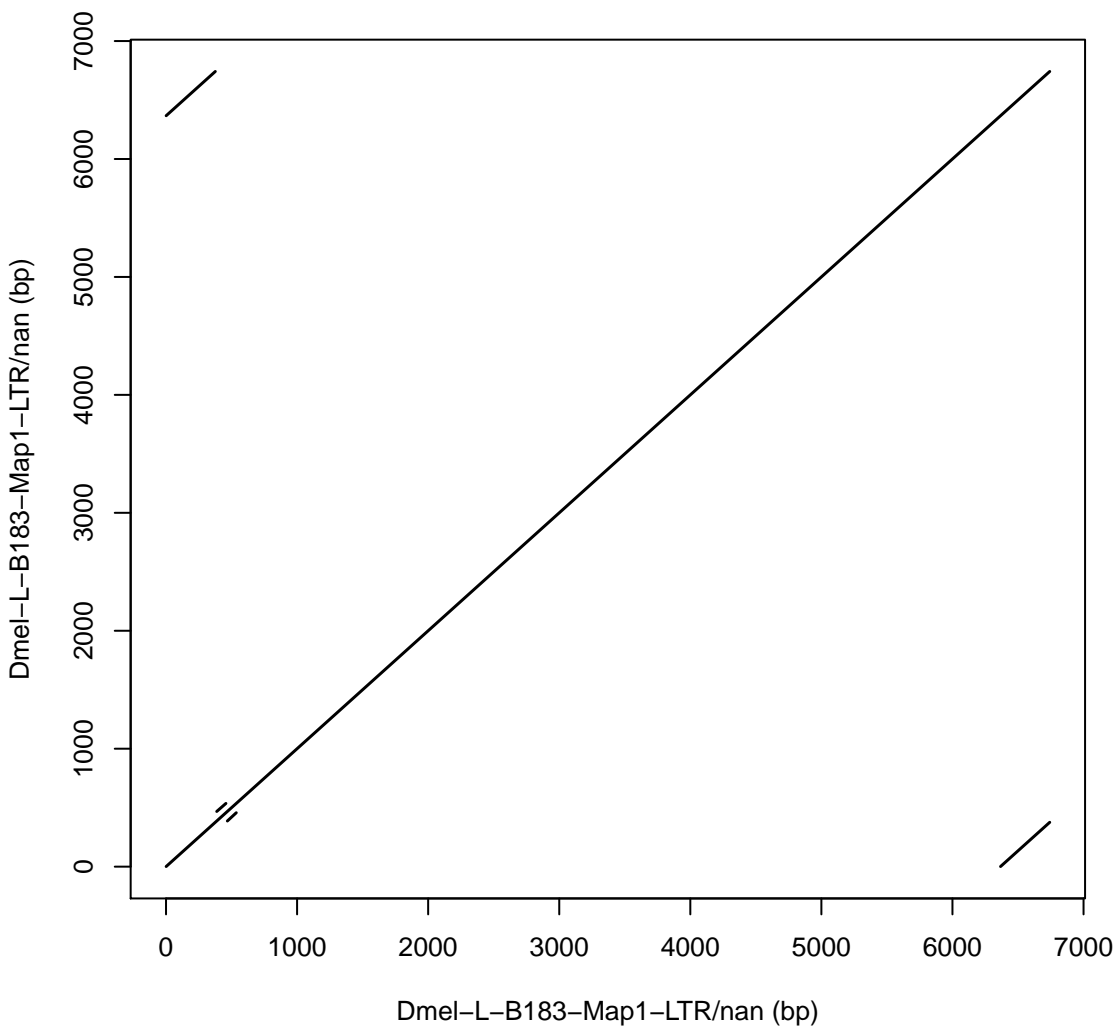
TE: Dmel-L-B183-Map1-LTR/nan  
consensus size: 6742bp; fragments: 295; full length: 1 (>=6067.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

