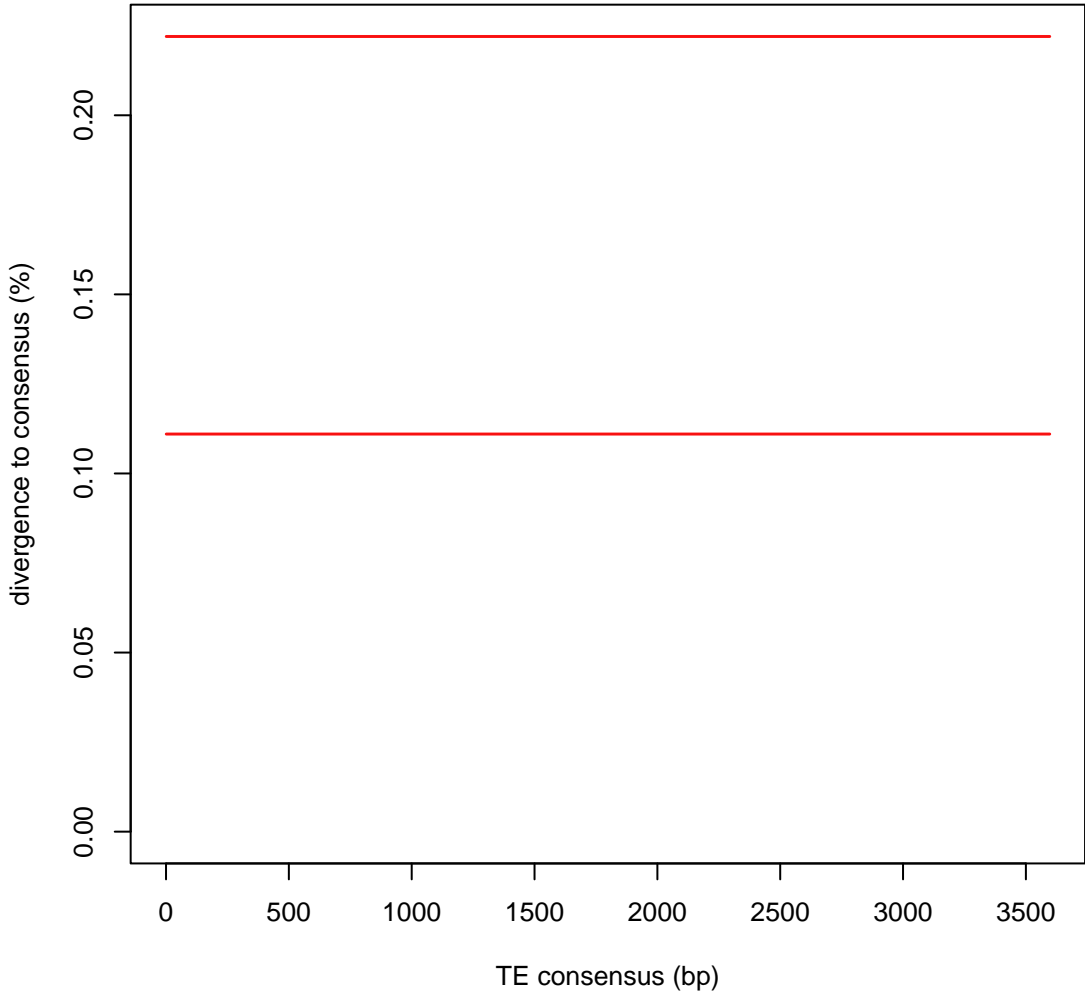
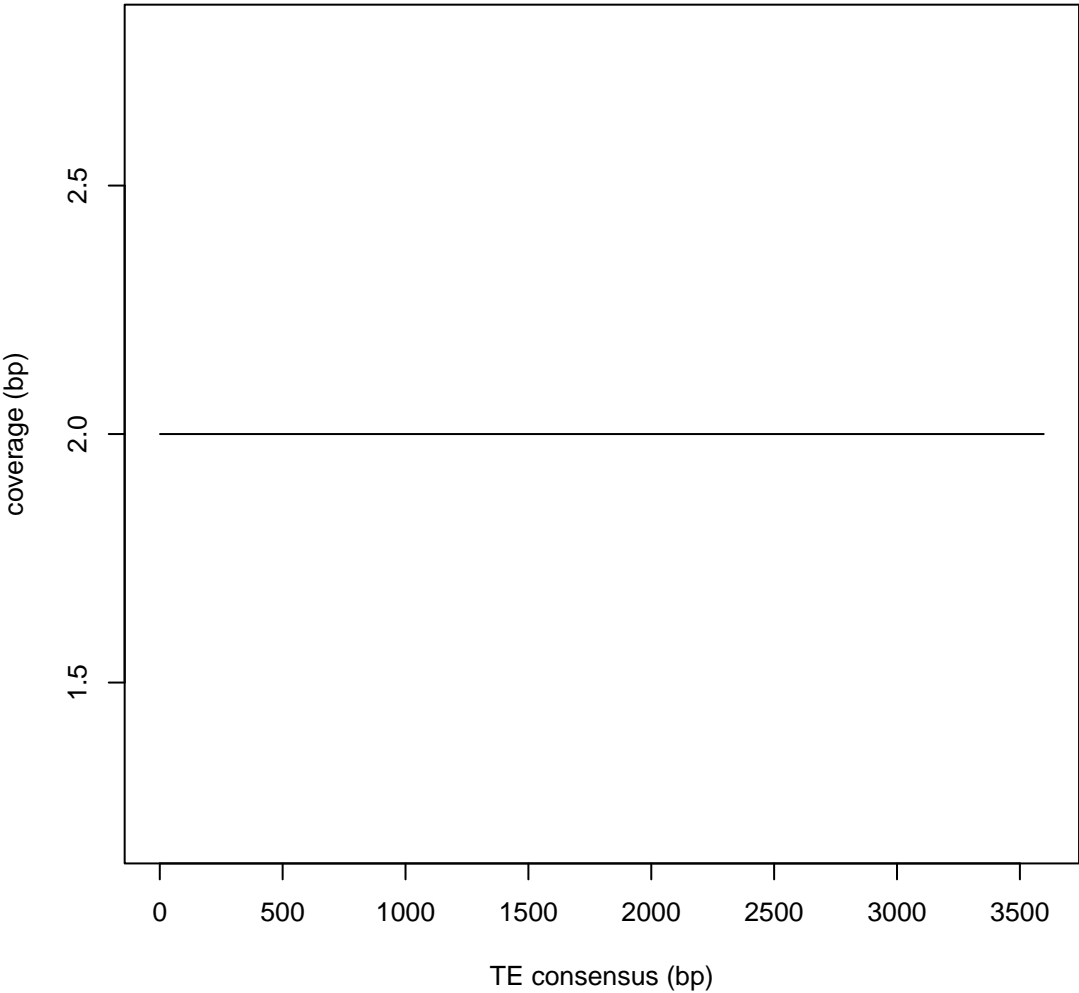


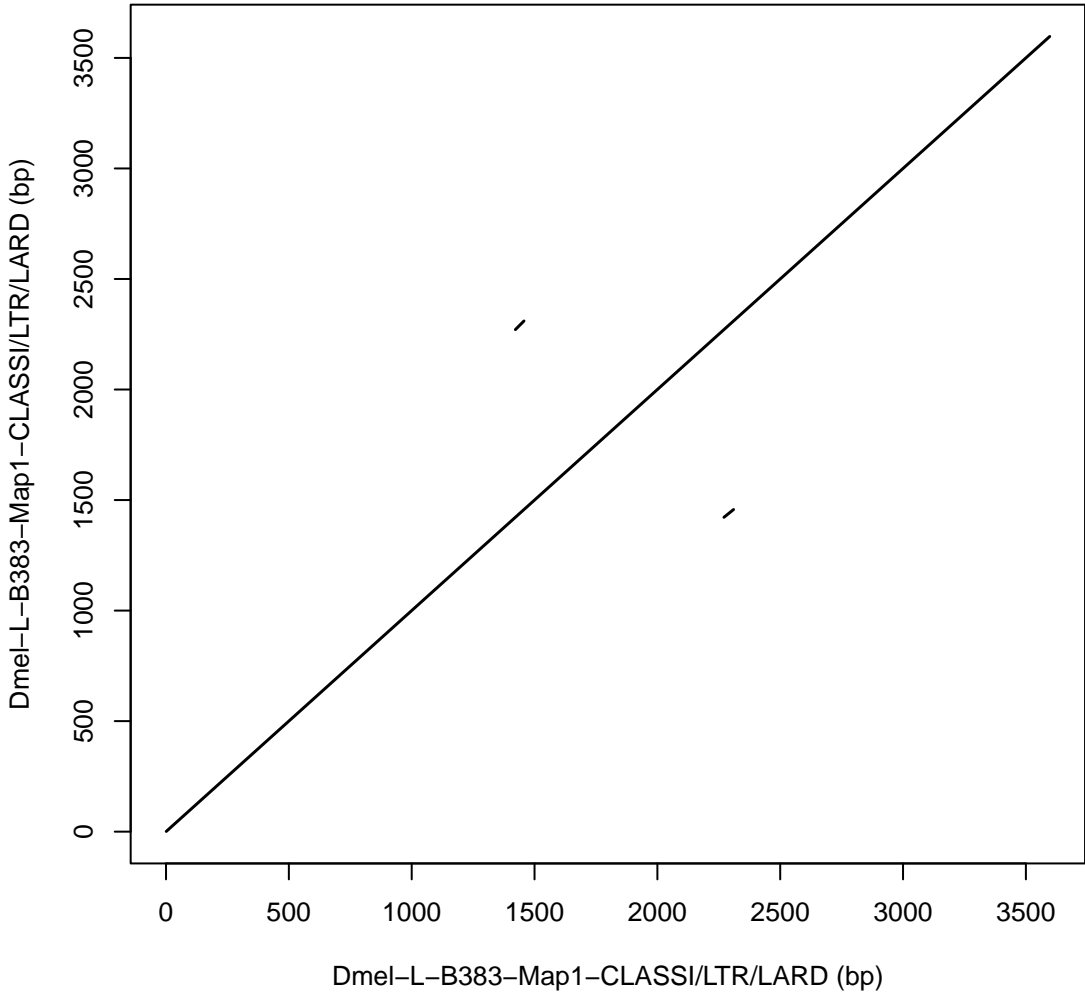
TE: Dmel-L-B383-Map1-CLASSI/LTR/LARD  
consensus size: 3597bp; fragments: 2; full length: 2 (>=3237.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

