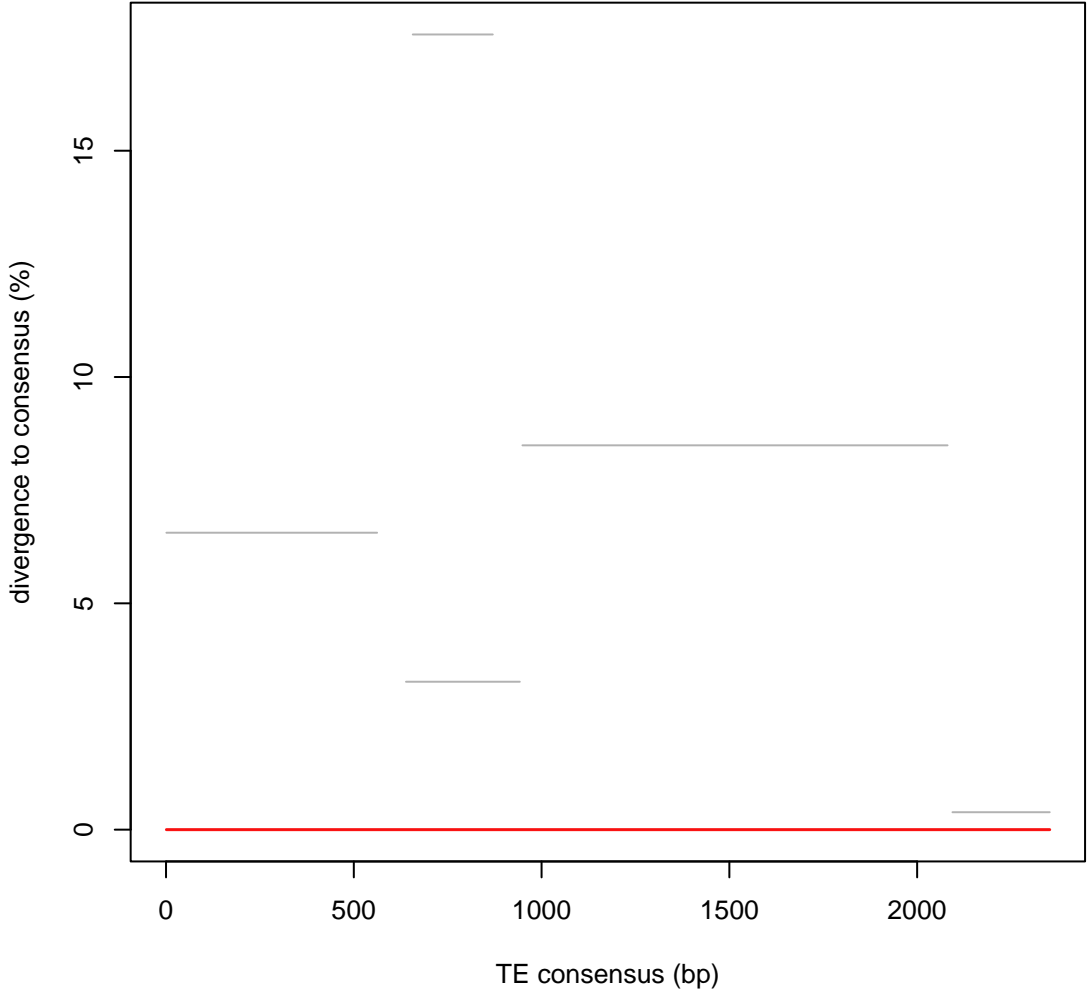
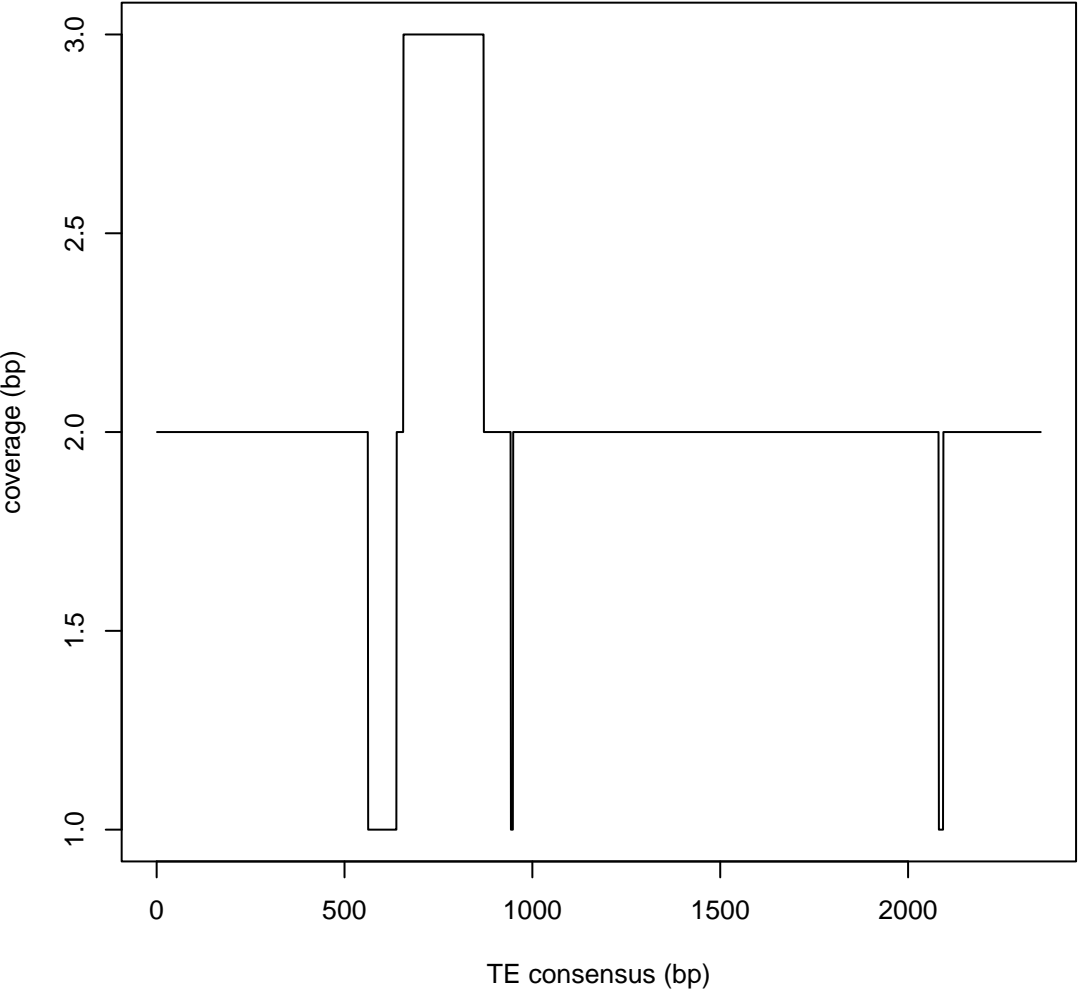


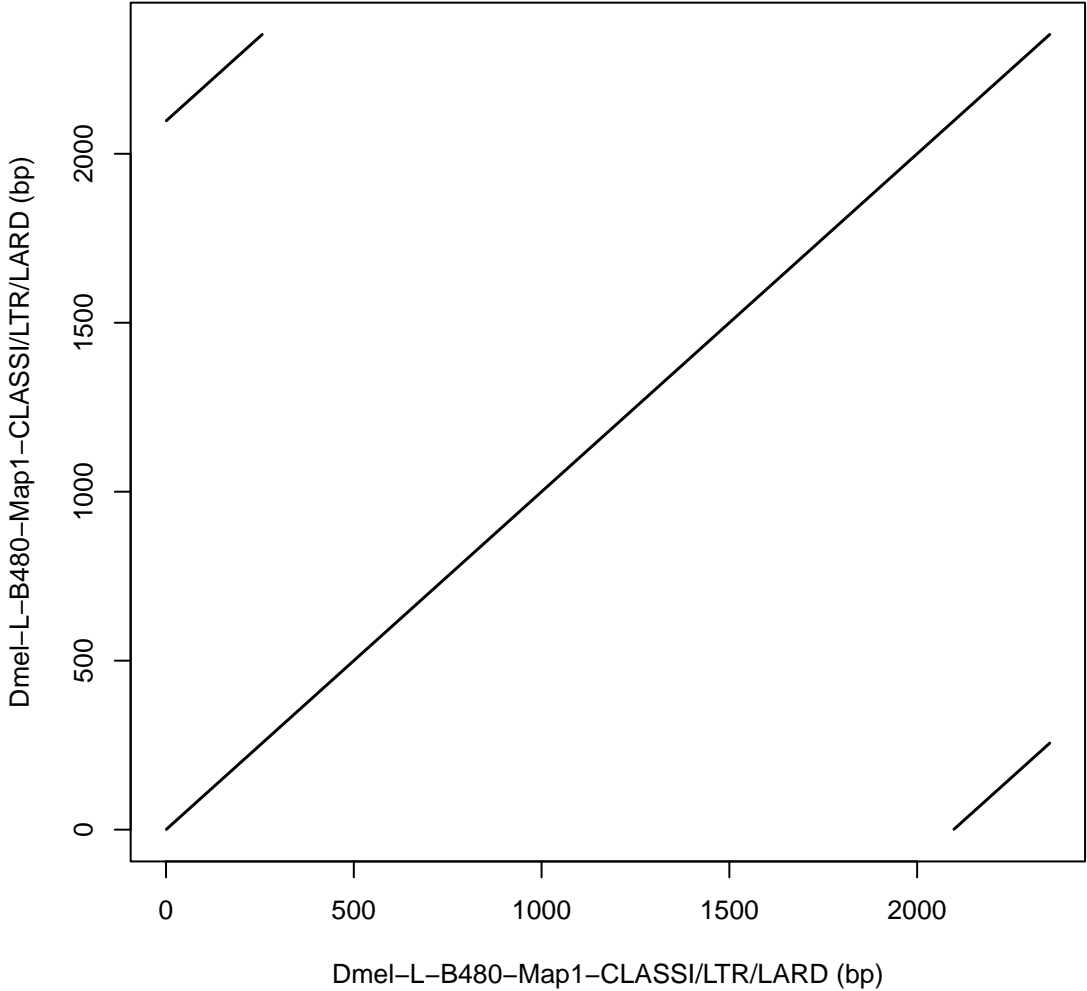
TE: Dmel-L-B480-Map1-CLASSI/LTR/LARD  
consensus size: 2353bp; fragments: 6; full length: 1 (>=2117.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

