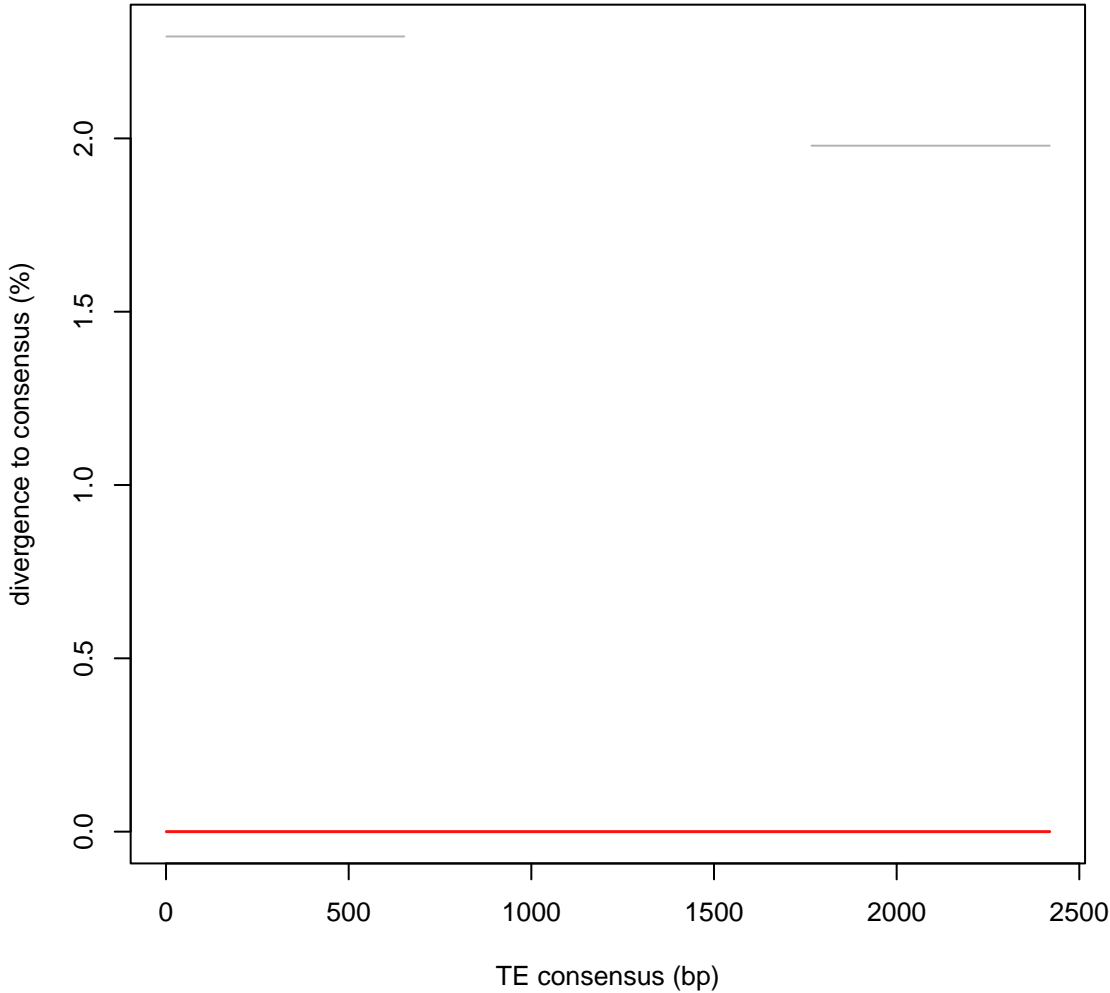
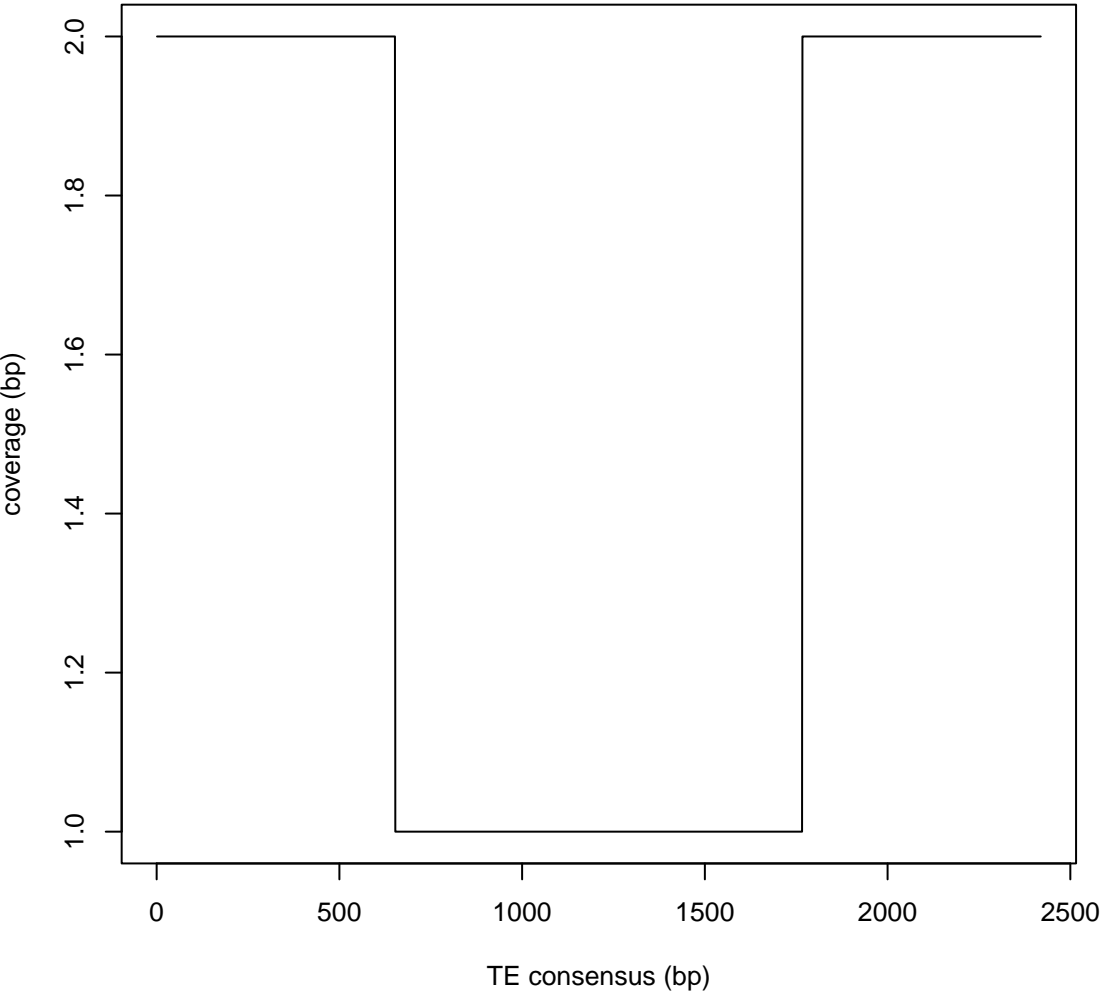


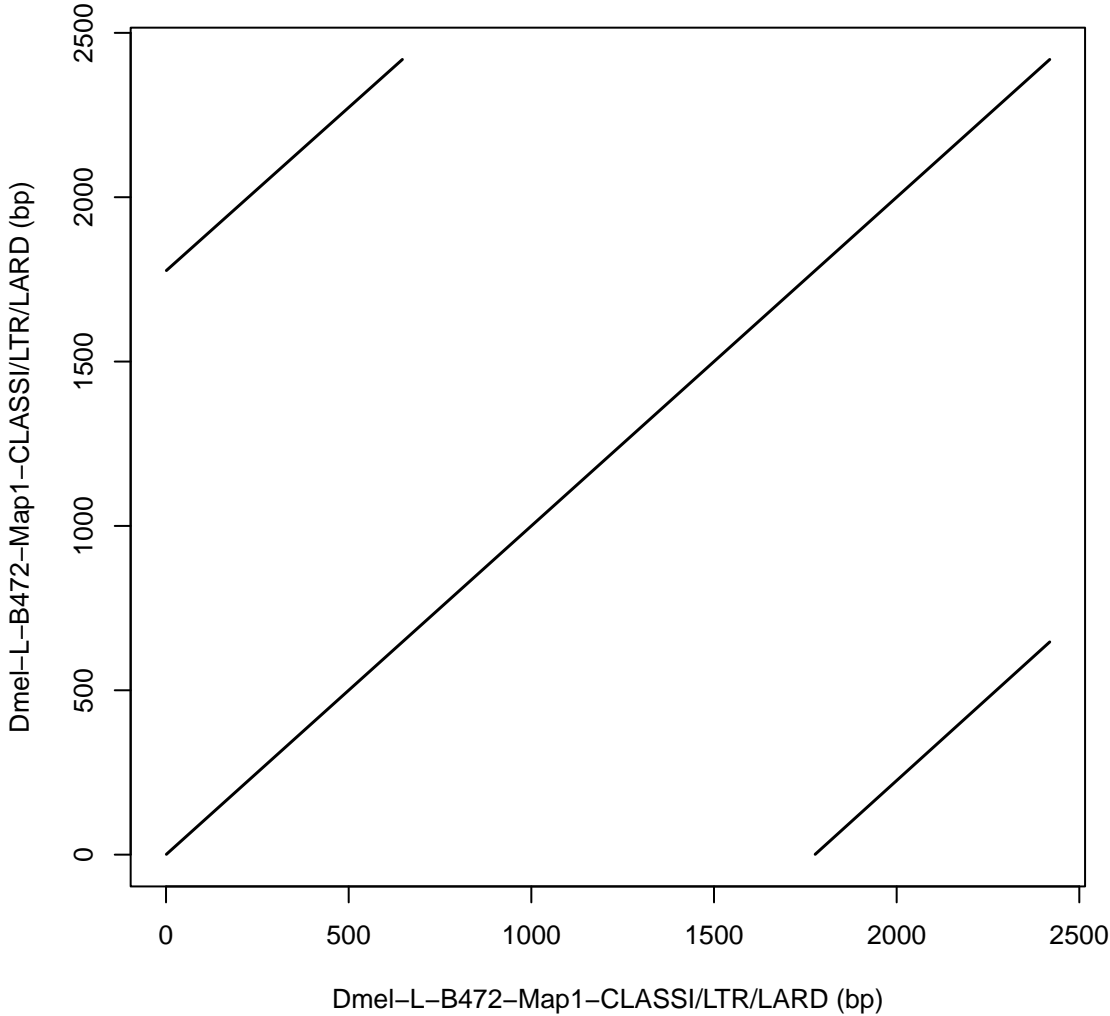
TE: Dmel-L-B472-Map1-CLASSI/LTR/LARD  
consensus size: 2419bp; fragments: 3; full length: 1 (>=2177.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

