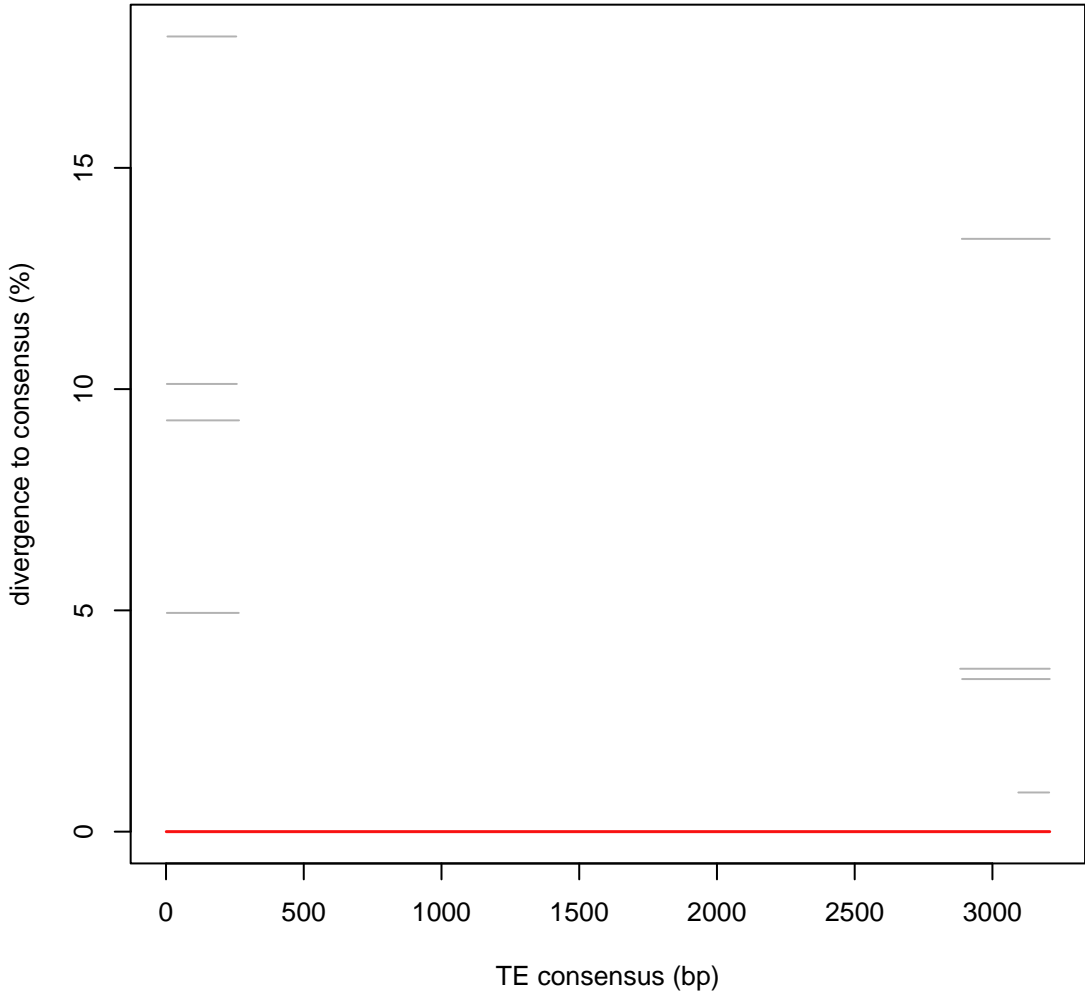
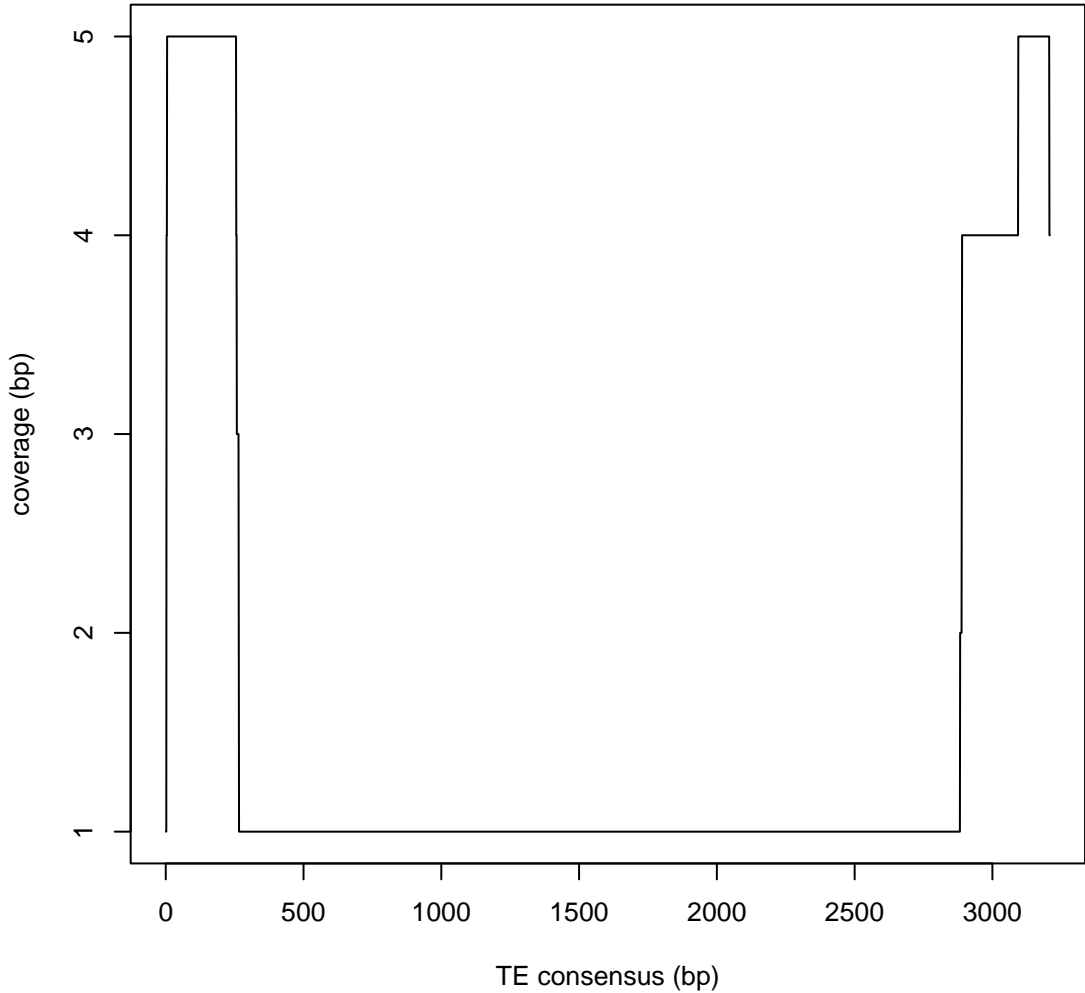


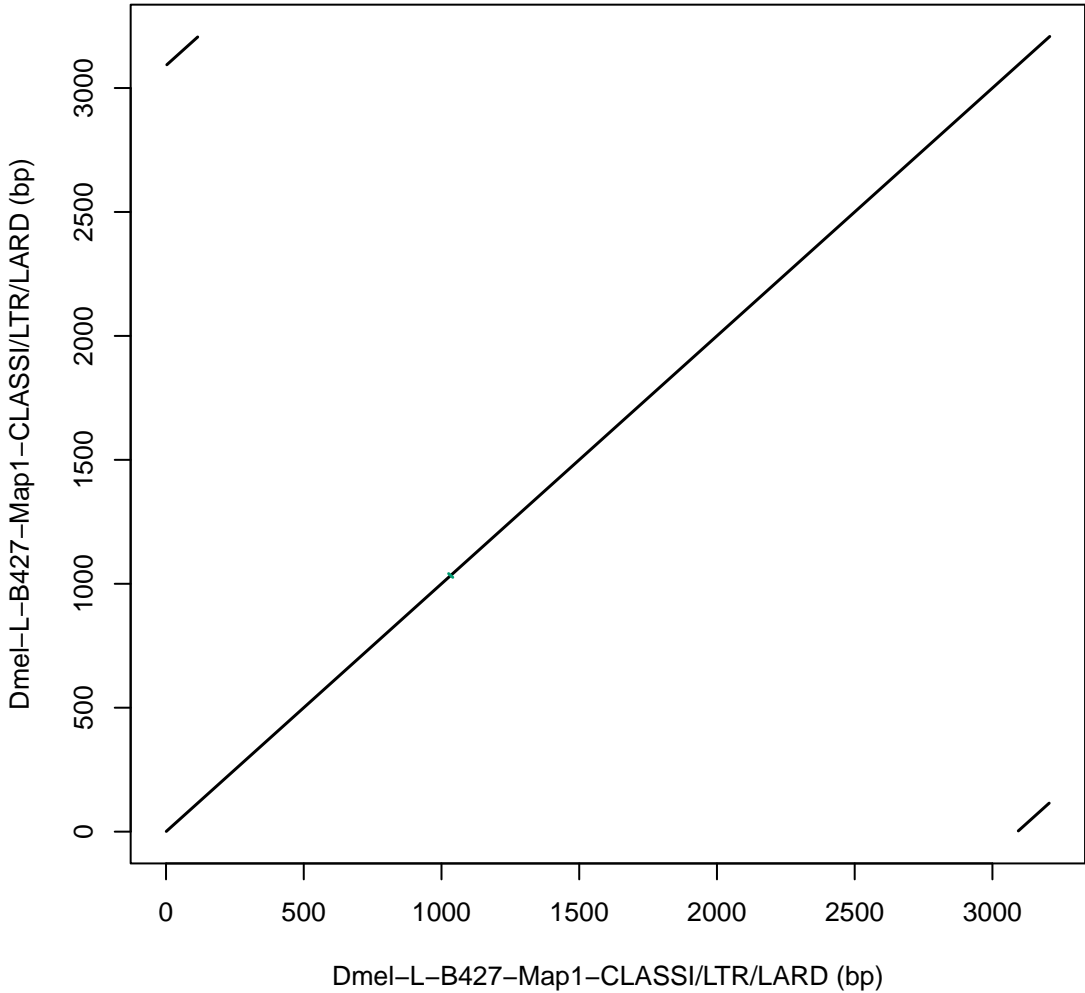
TE: Dmel-L-B427-Map1-CLASSI/LTR/LARD  
consensus size: 3208bp; fragments: 9; full length: 1 (>=2887.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

