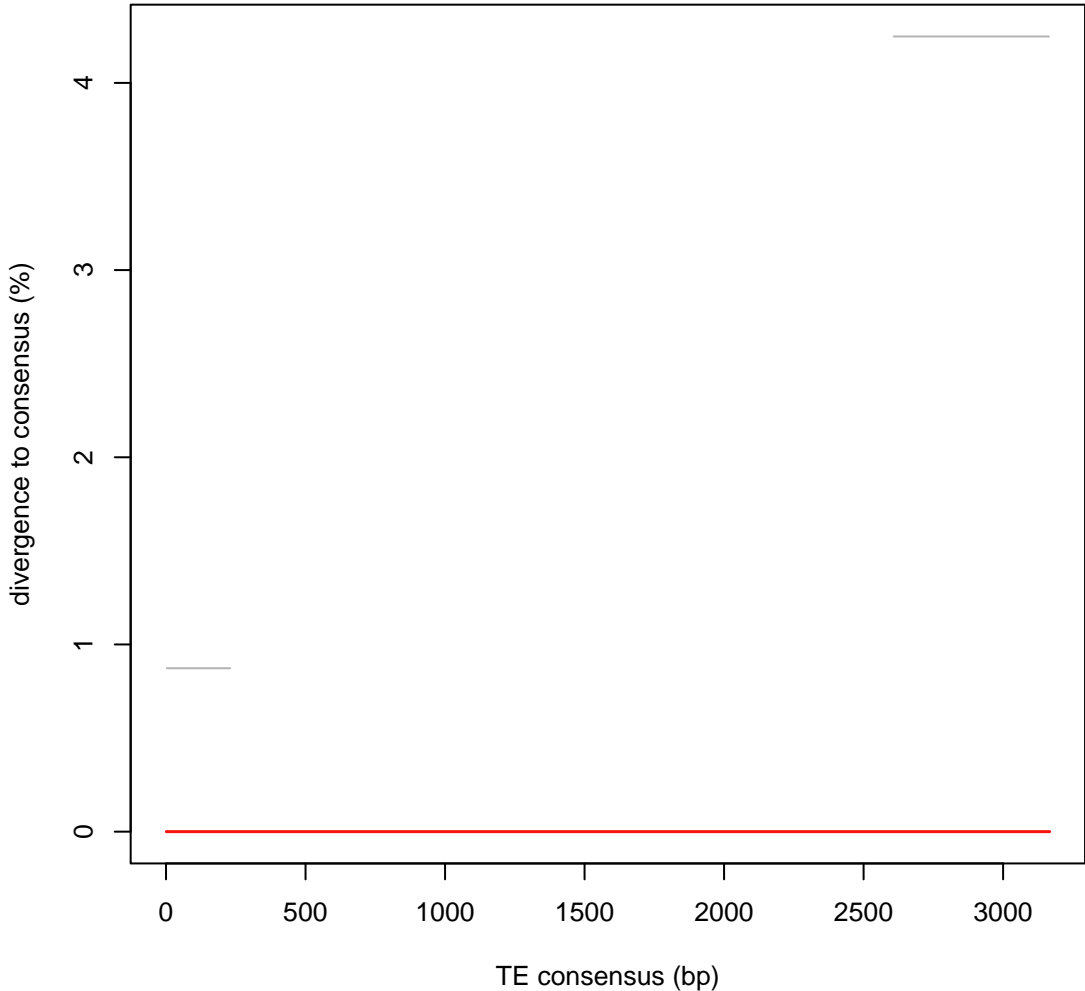
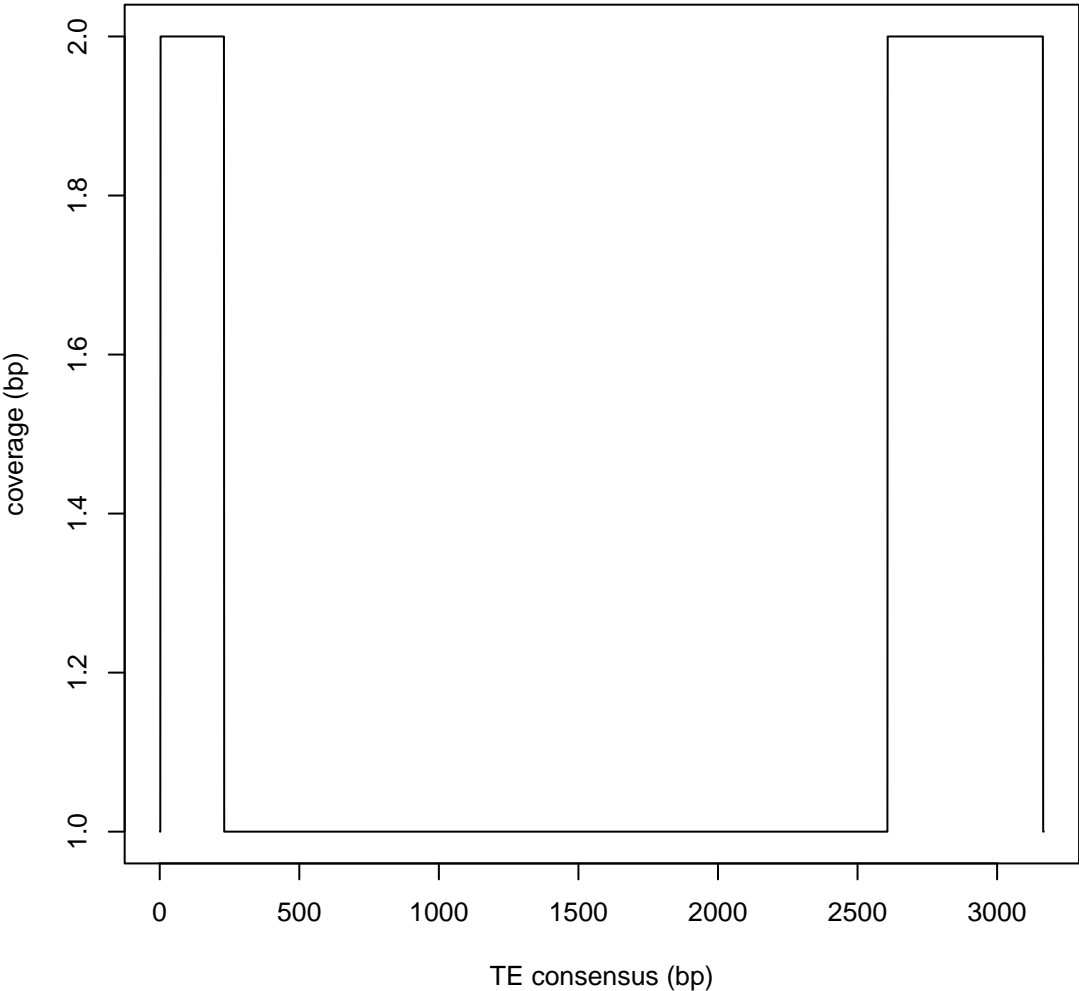


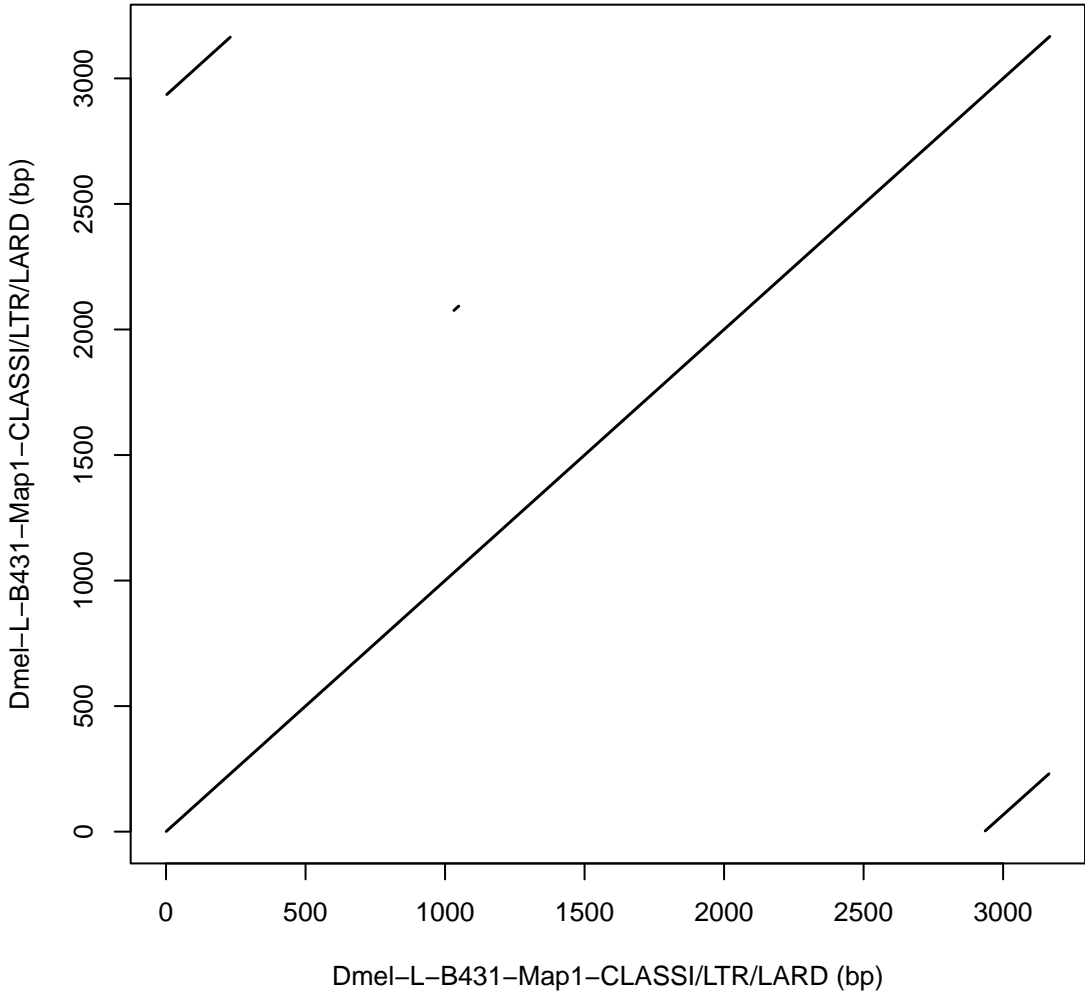
TE: Dmel-L-B431-Map1-CLASSI/LTR/LARD
consensus size: 3167bp; fragments: 3; full length: 1 (>=2850.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

