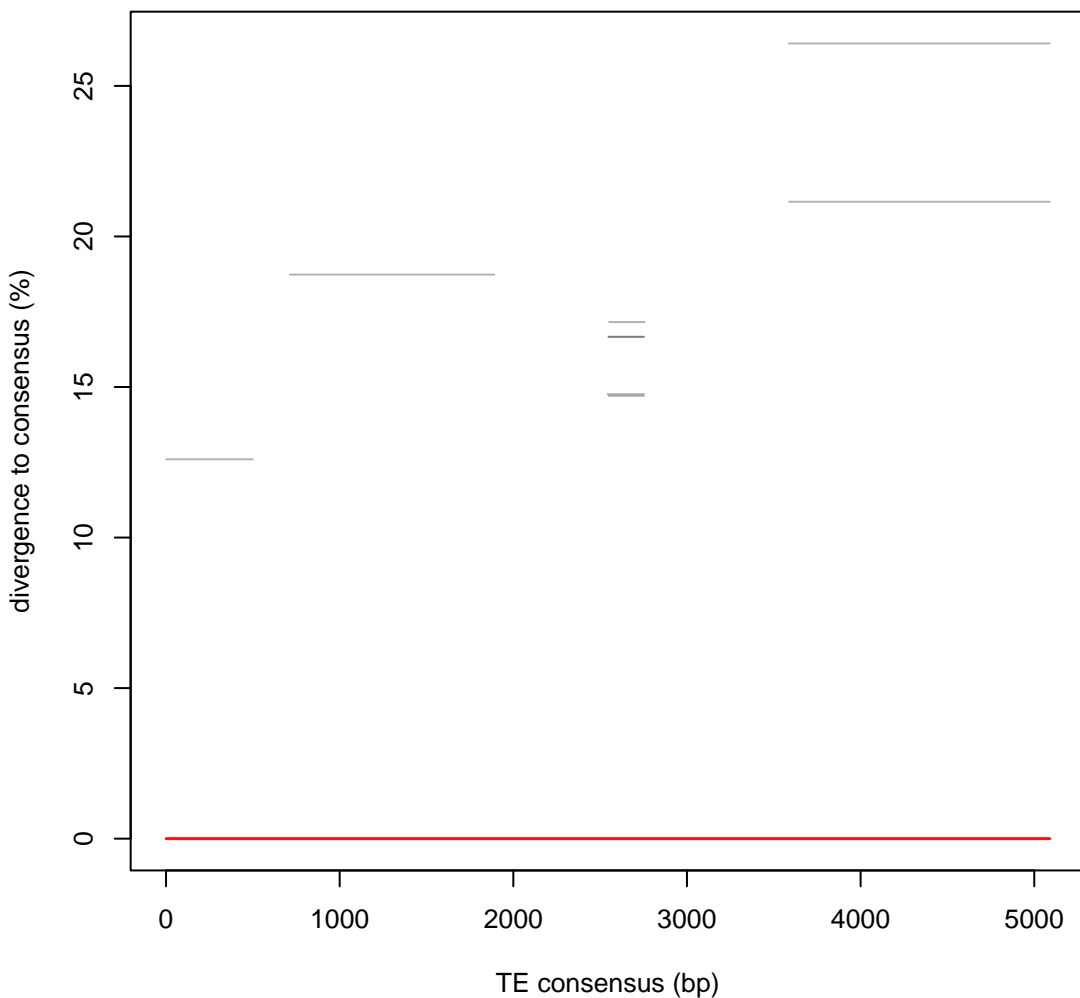
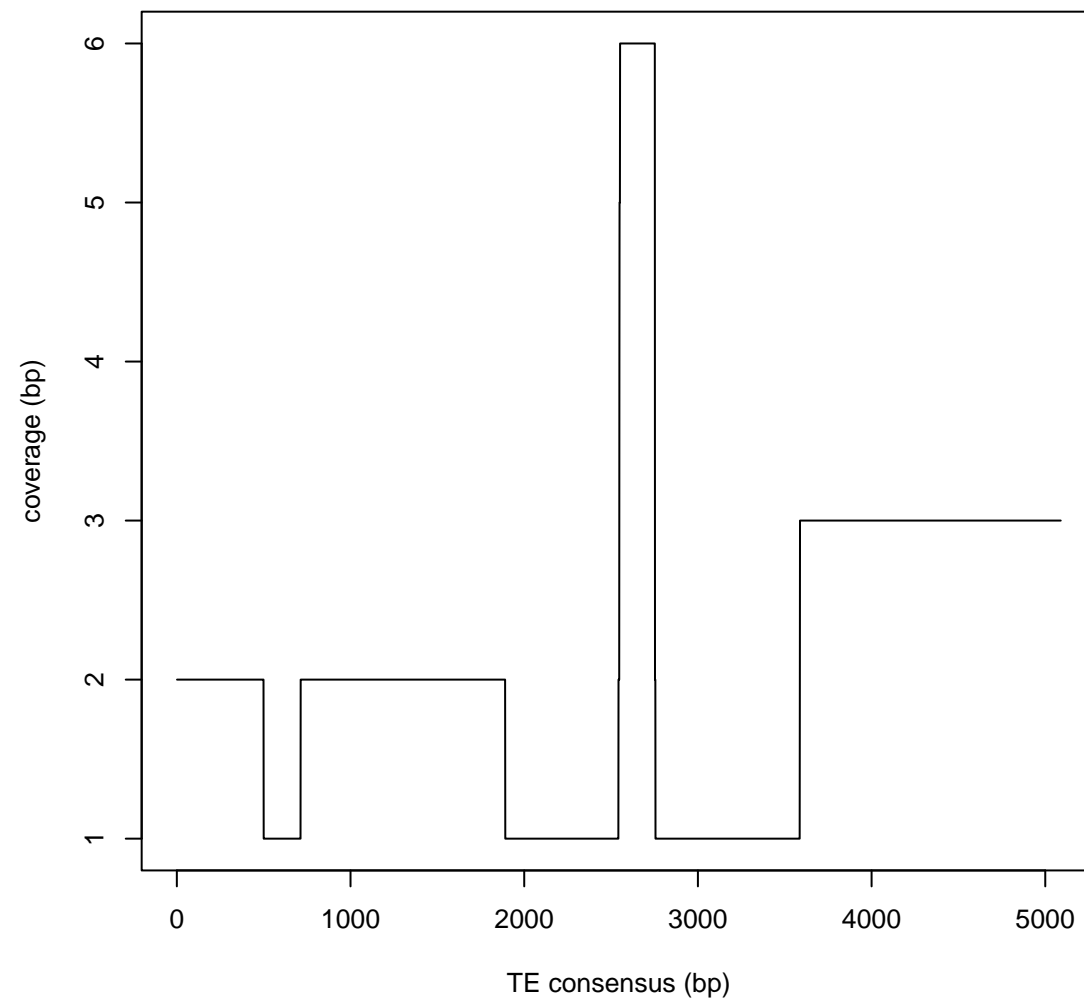


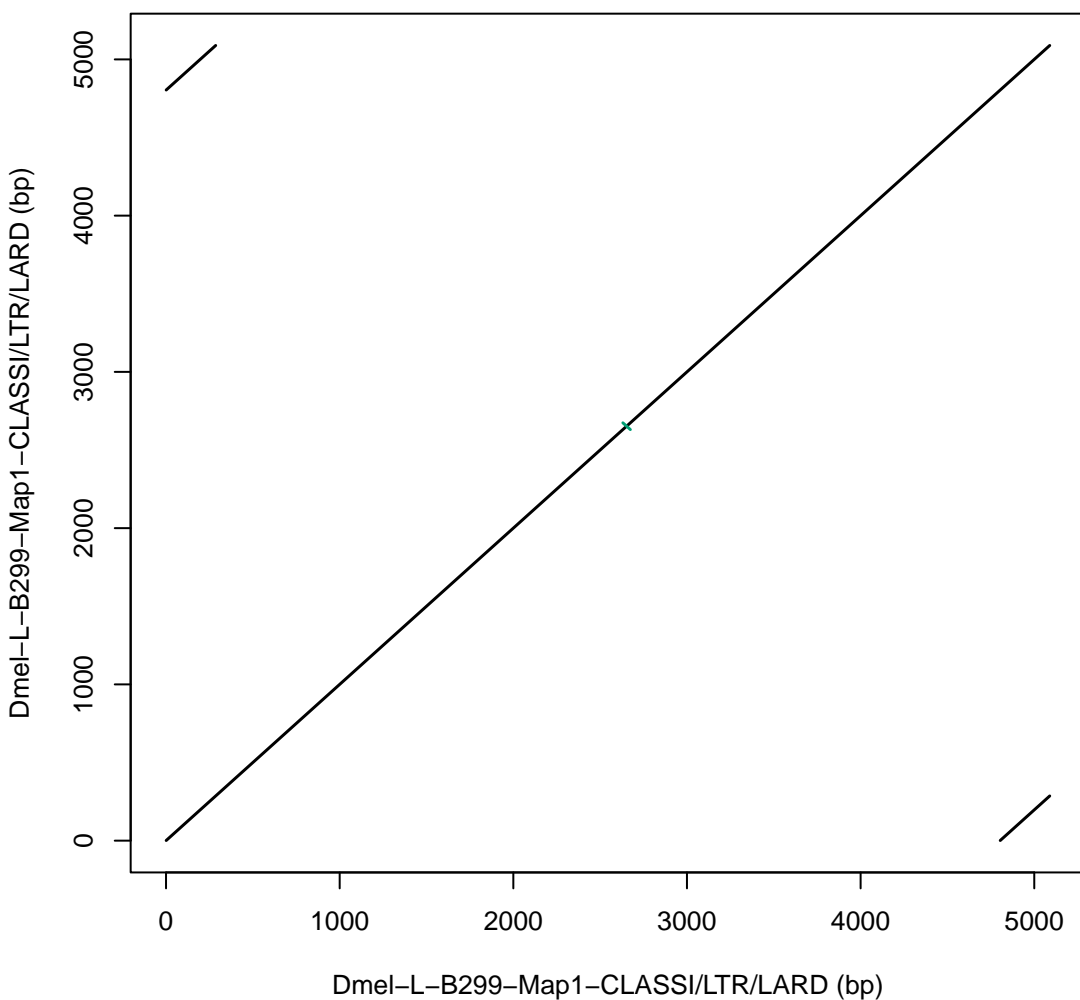
TE: Dmel-L-B299-Map1-CLASSI/LTR/LARD
consensus size: 5089bp; fragments: 10; full length: 1 (≥ 4580.1 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

