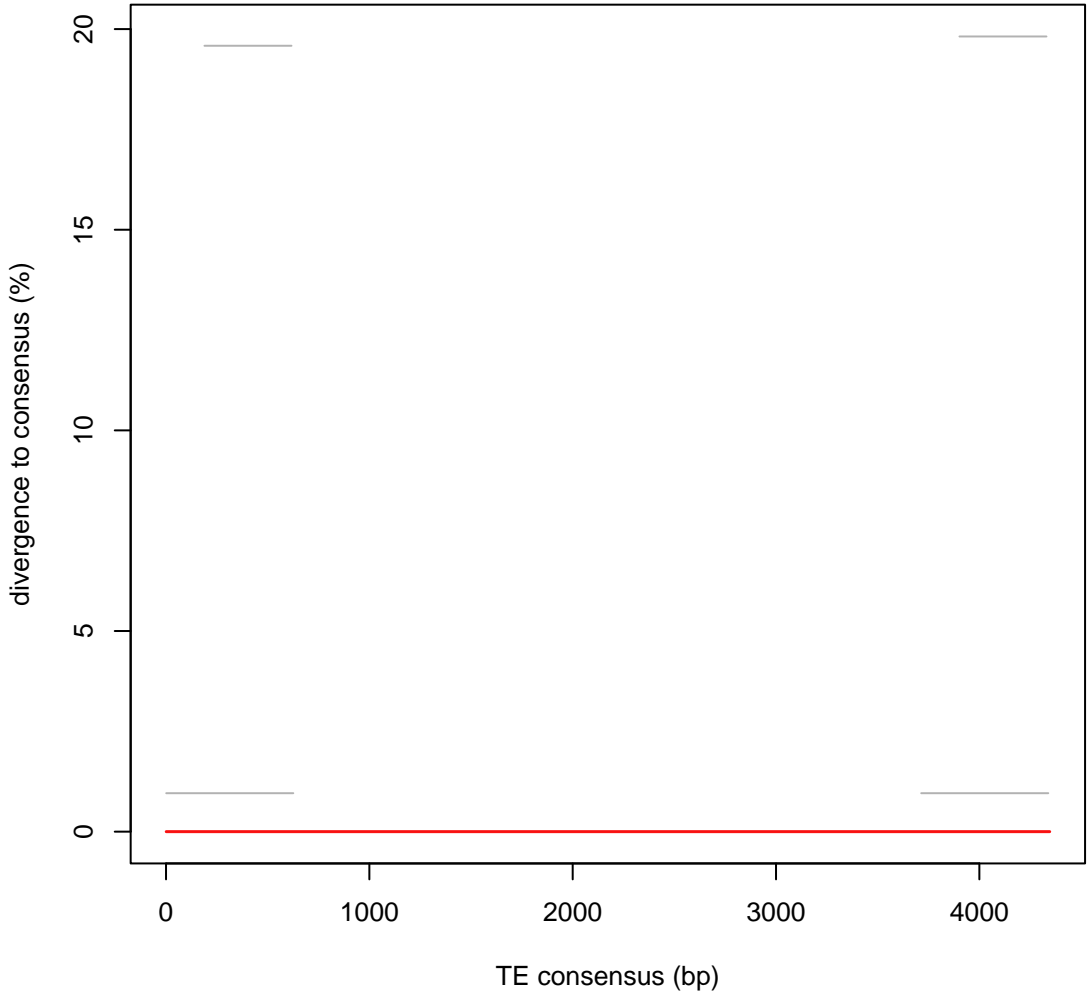
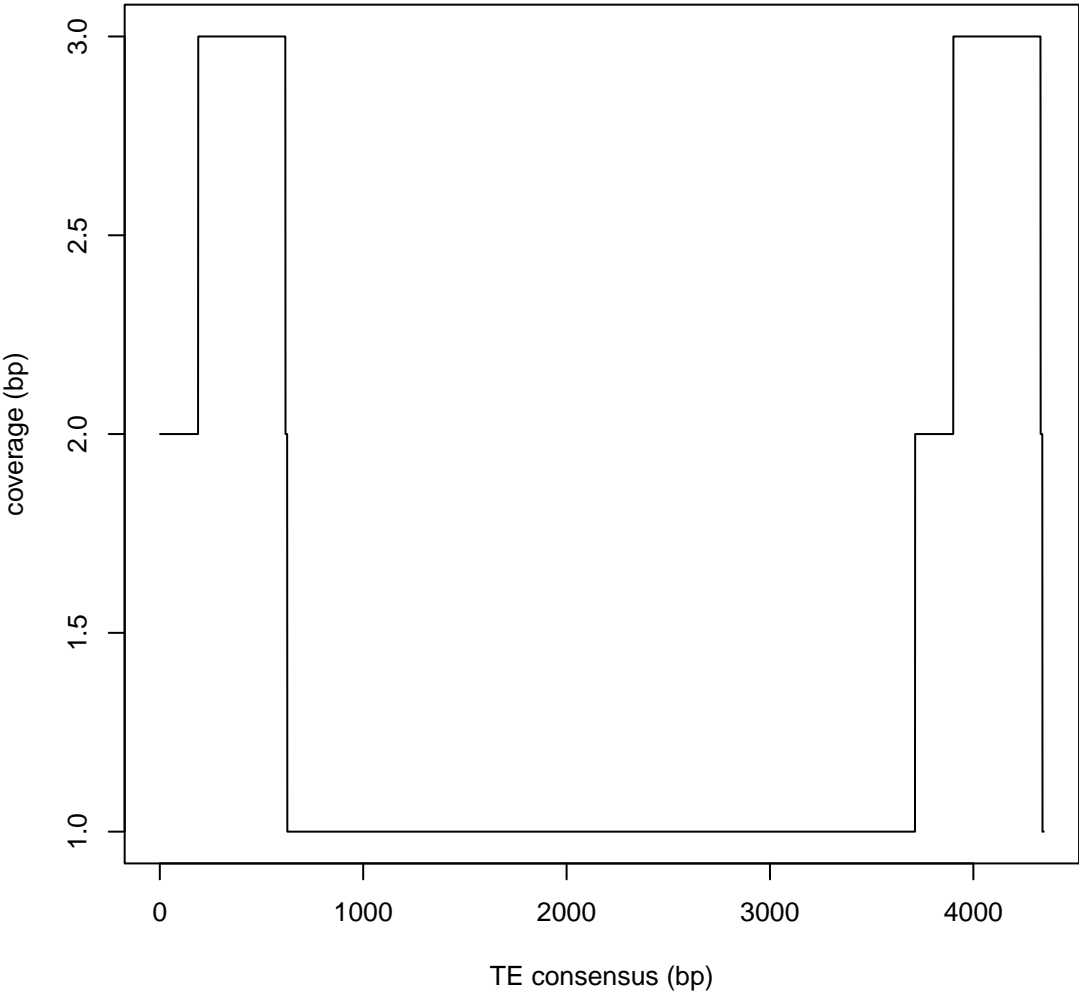


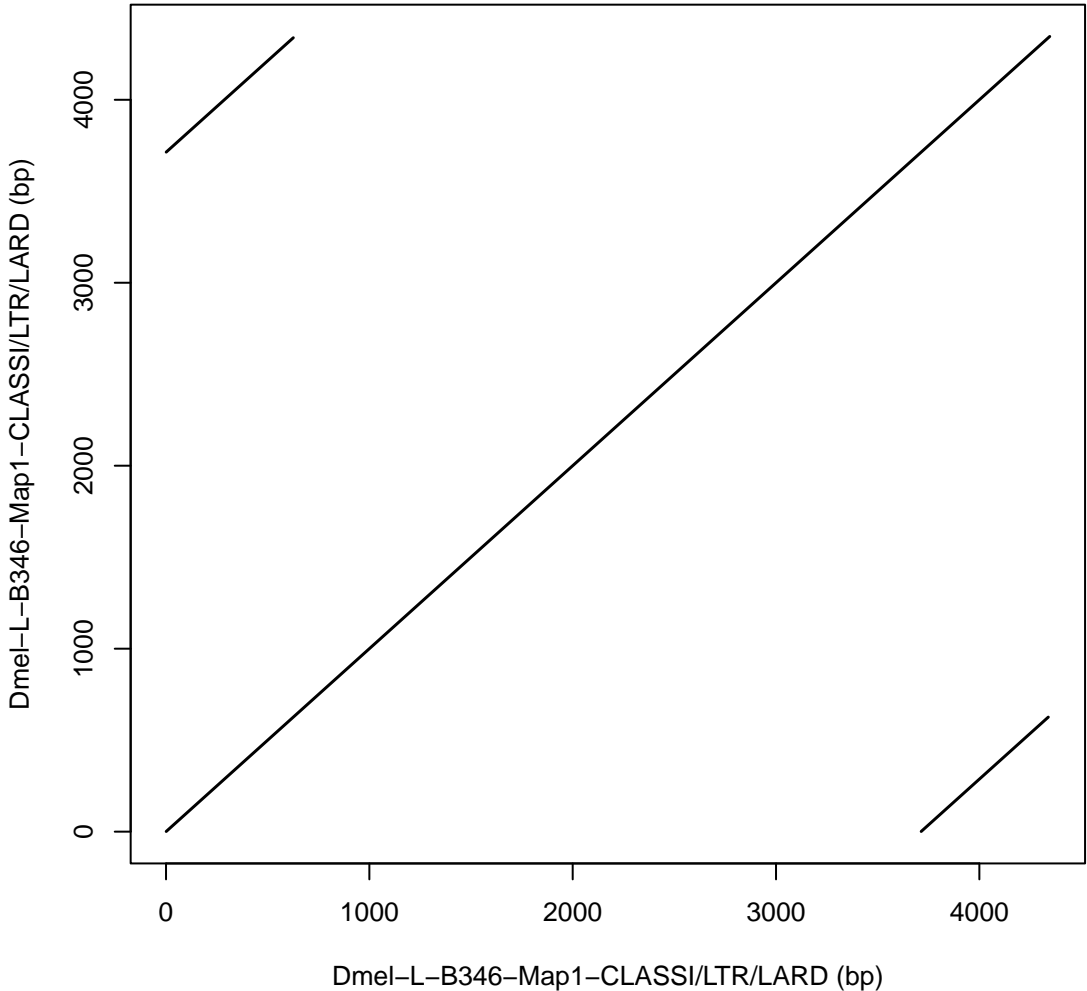
TE: Dmel-L-B346-Map1-CLASSI/LTR/LARD
consensus size: 4346bp; fragments: 5; full length: 1 (>=3911.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

