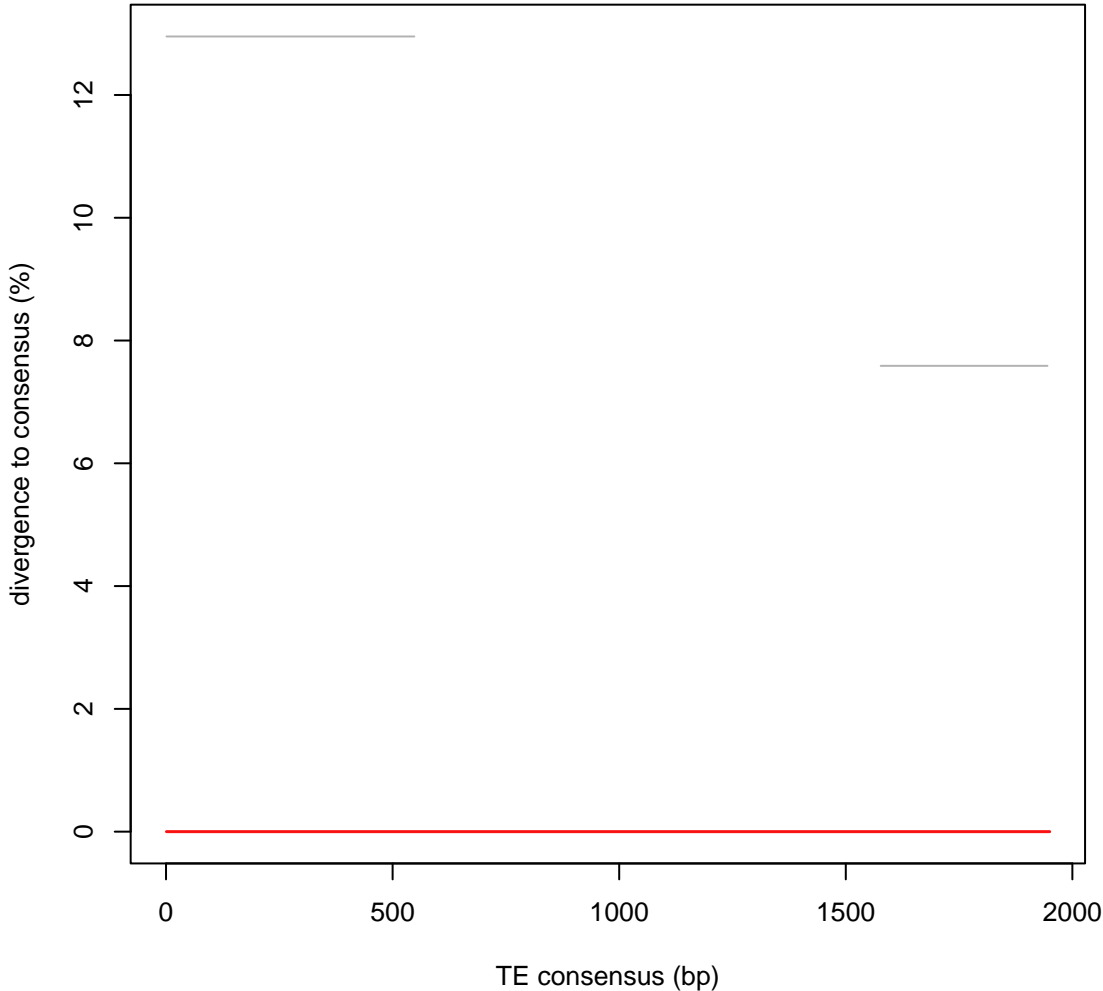
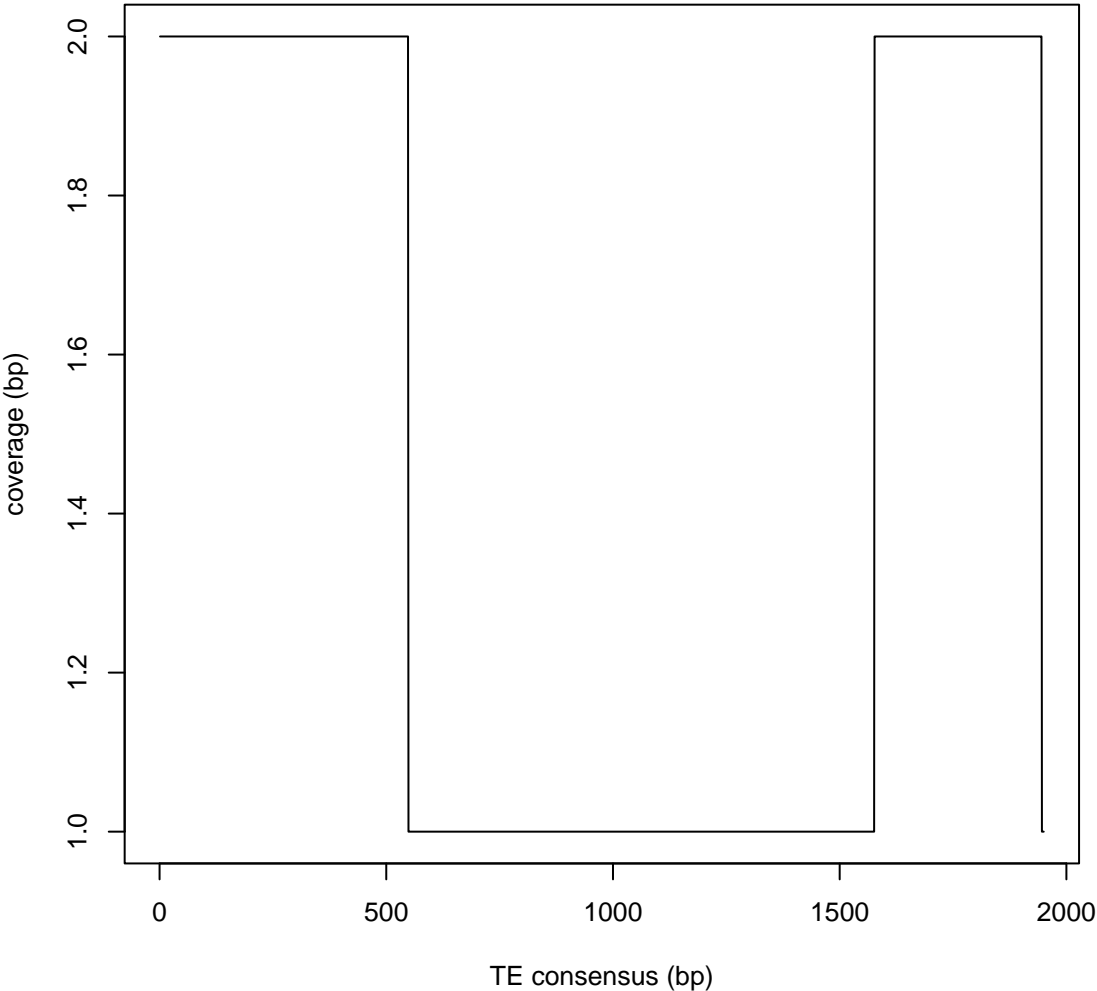


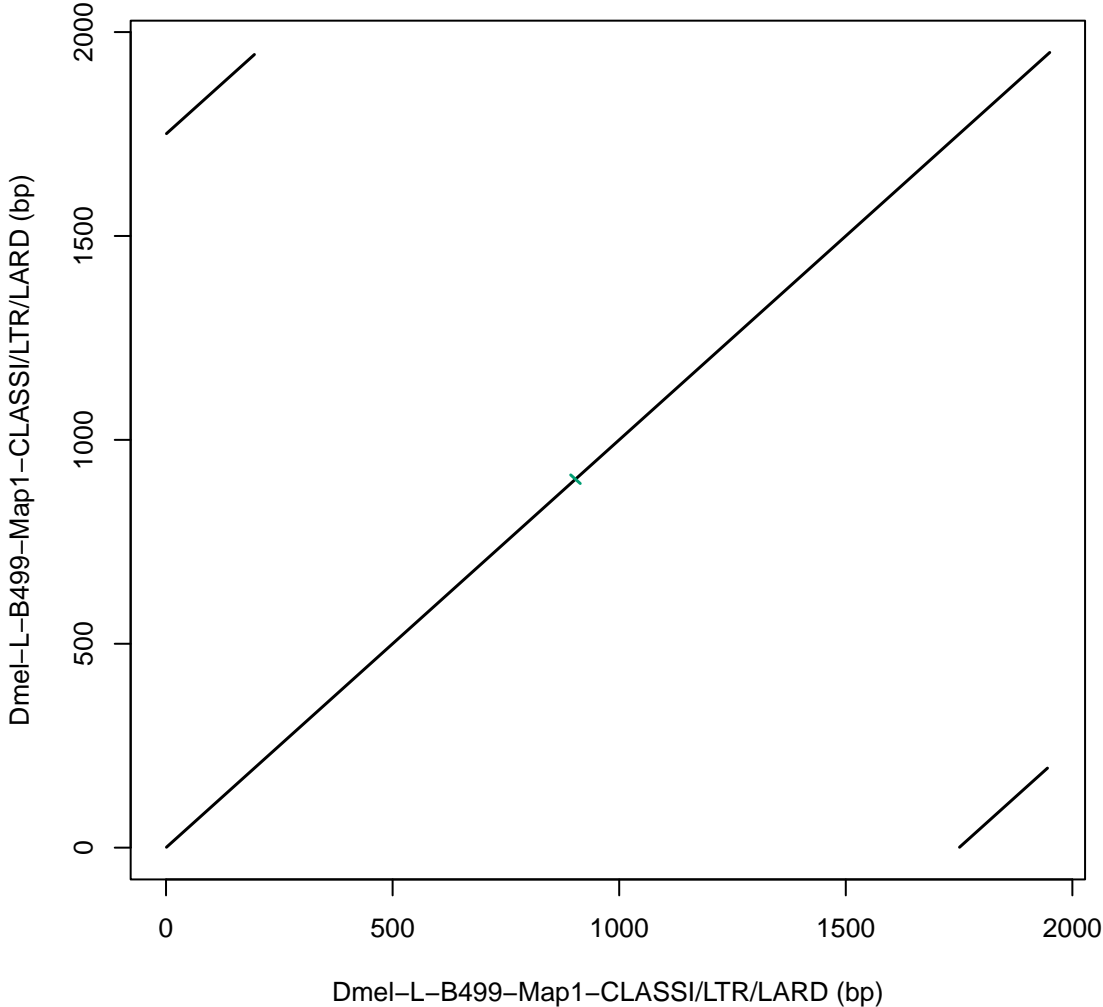
TE: Dmel-L-B499-Map1-CLASSI/LTR/LARD
consensus size: 1950bp; fragments: 3; full length: 1 (>=1755bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

