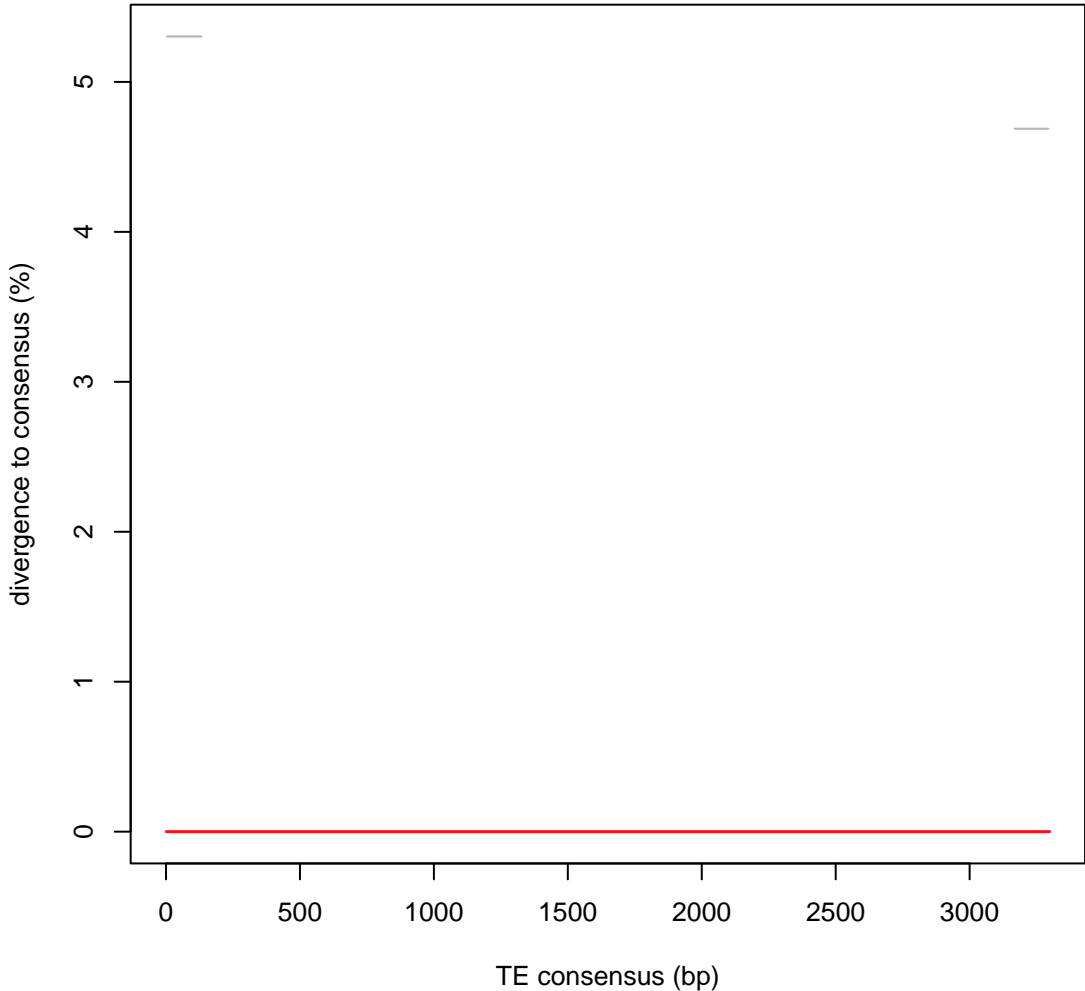
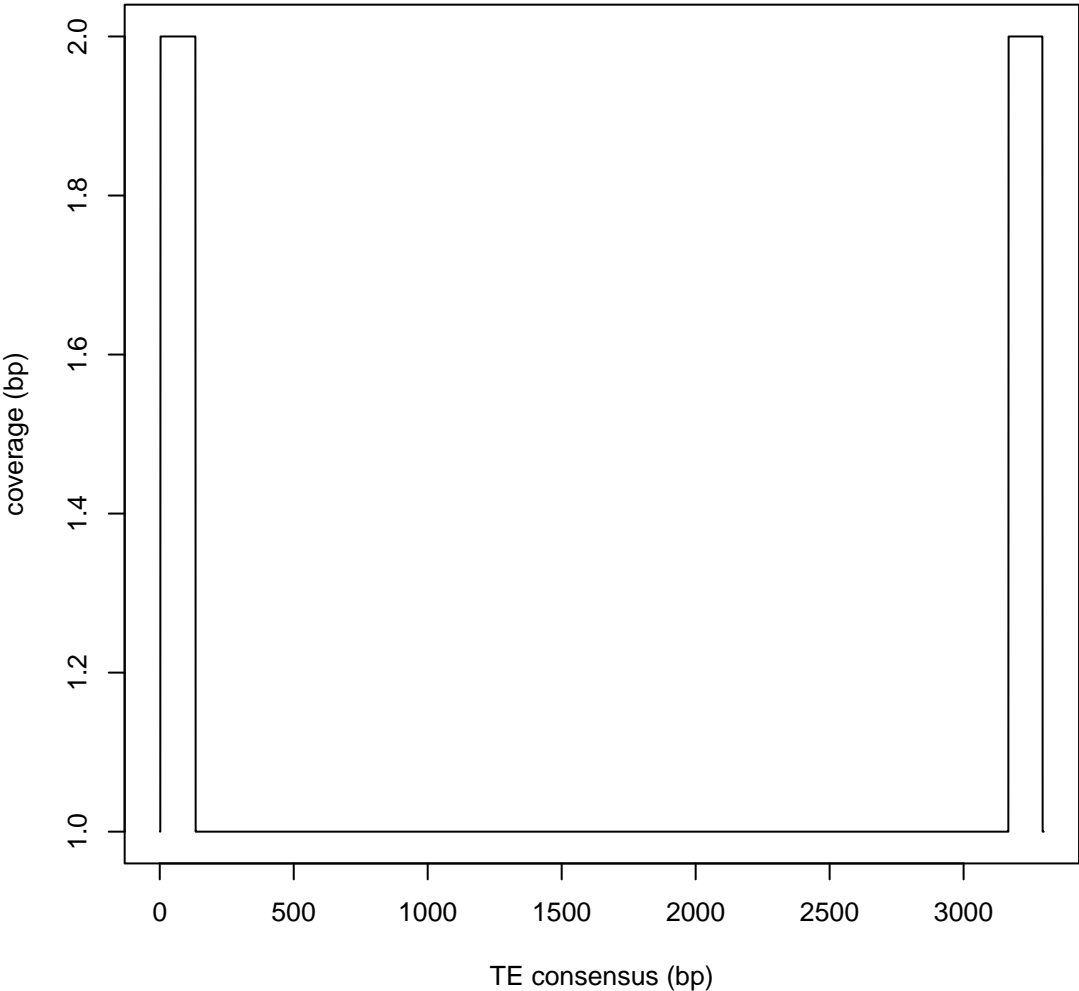


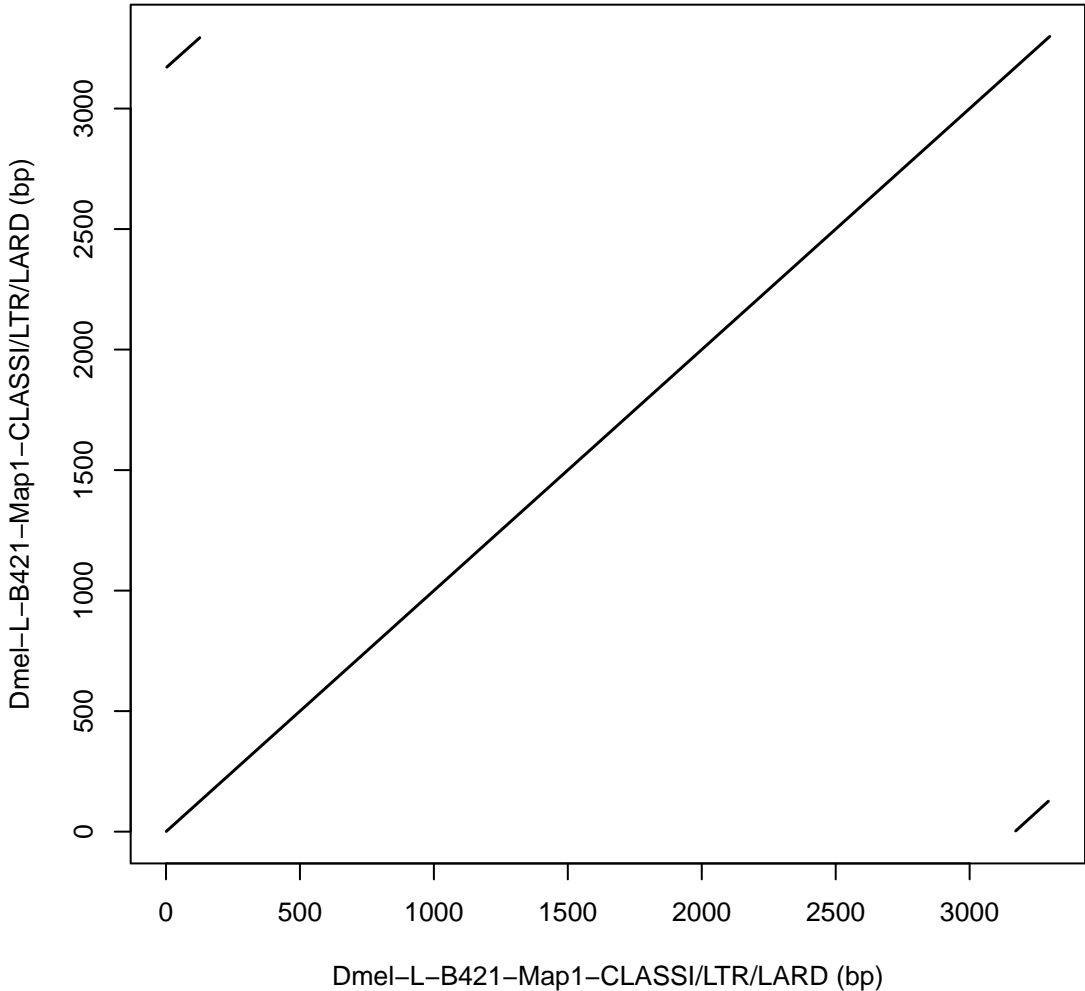
TE: Dmel-L-B421-Map1-CLASSI/LTR/LARD
consensus size: 3299bp; fragments: 3; full length: 1 (>=2969.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

