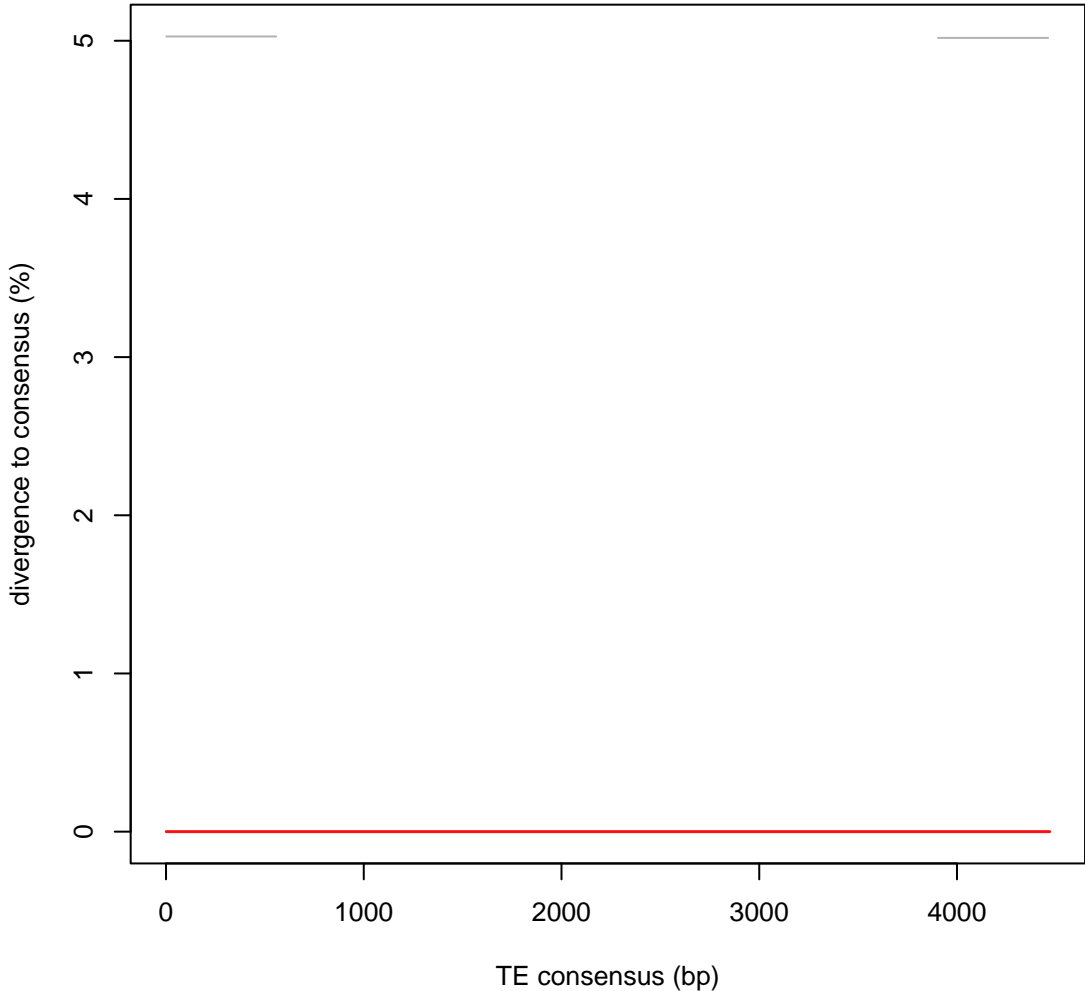
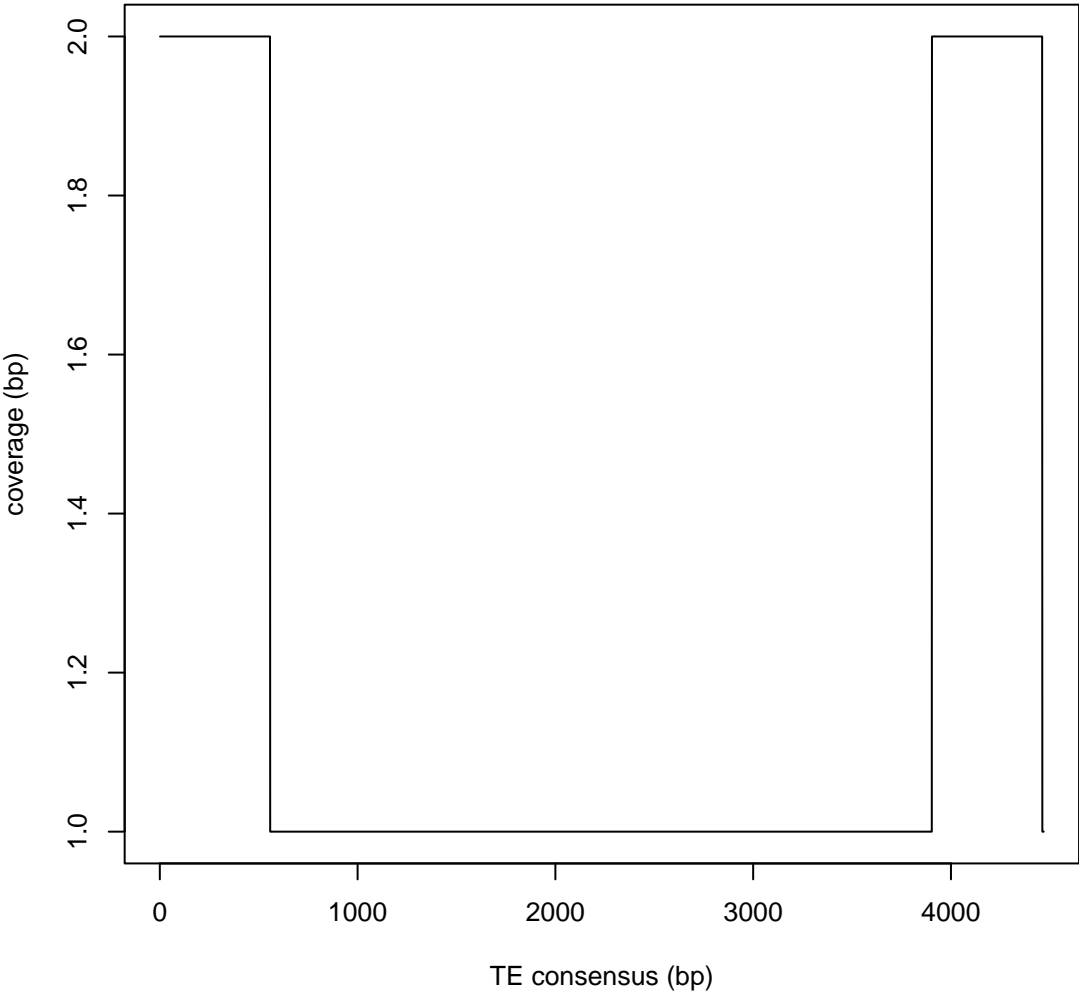


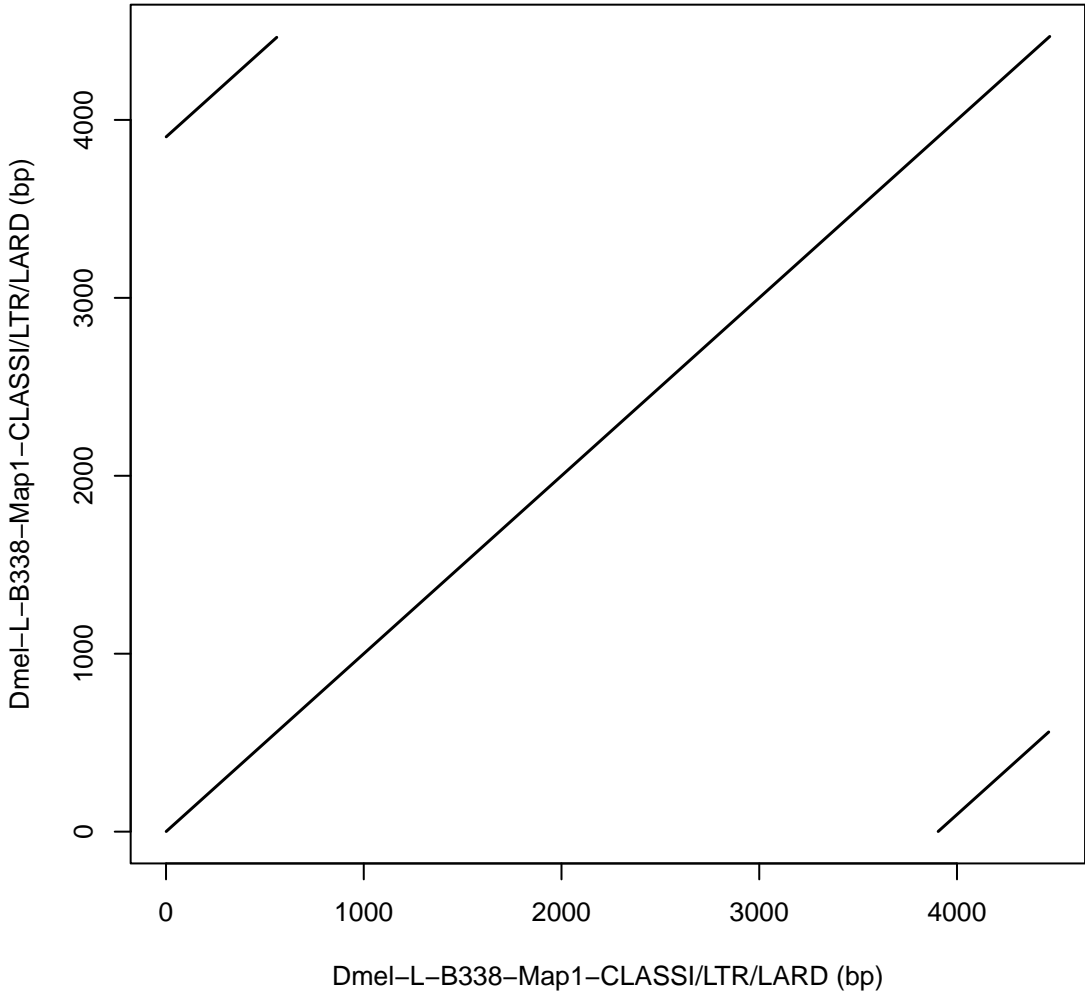
TE: Dmel-L-B338-Map1-CLASSI/LTR/LARD
consensus size: 4469bp; fragments: 3; full length: 1 (>=4022.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

