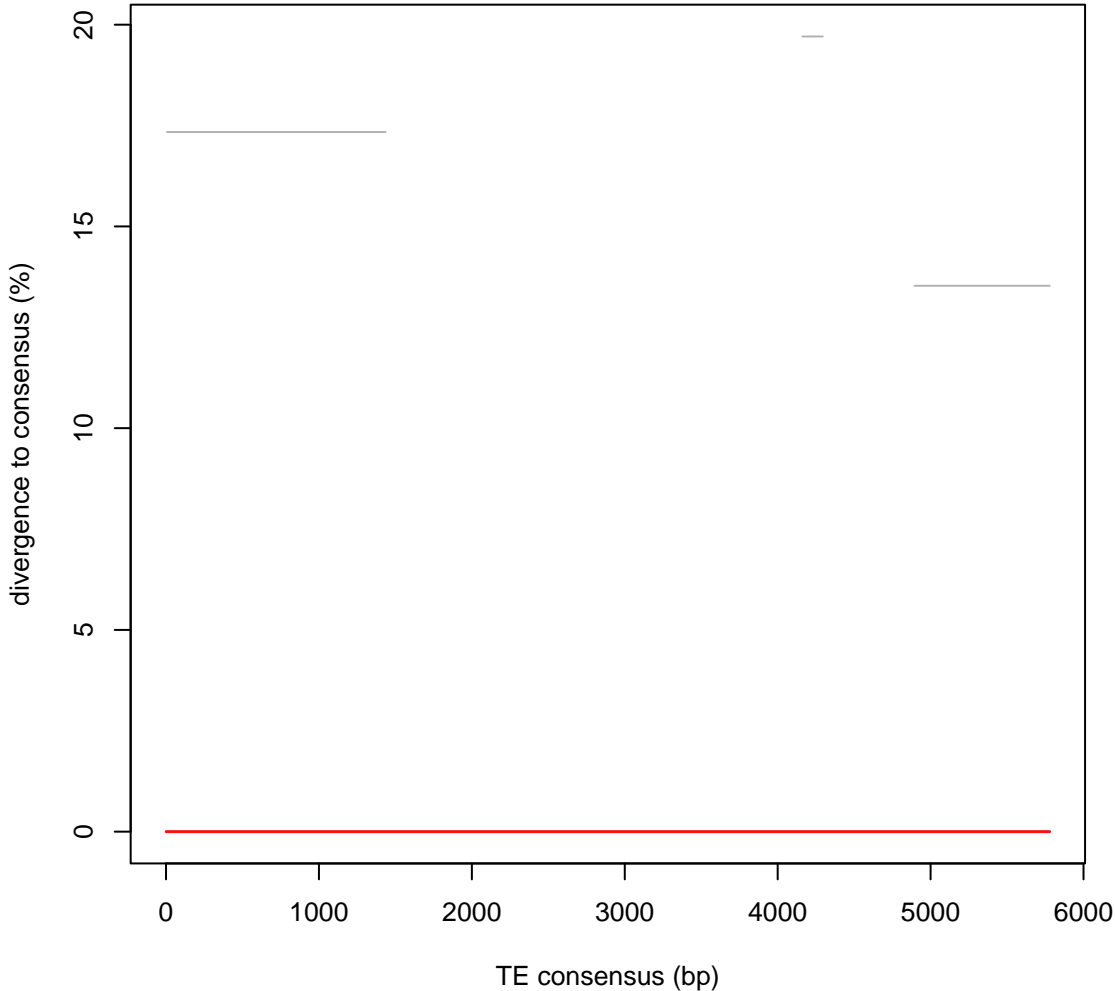
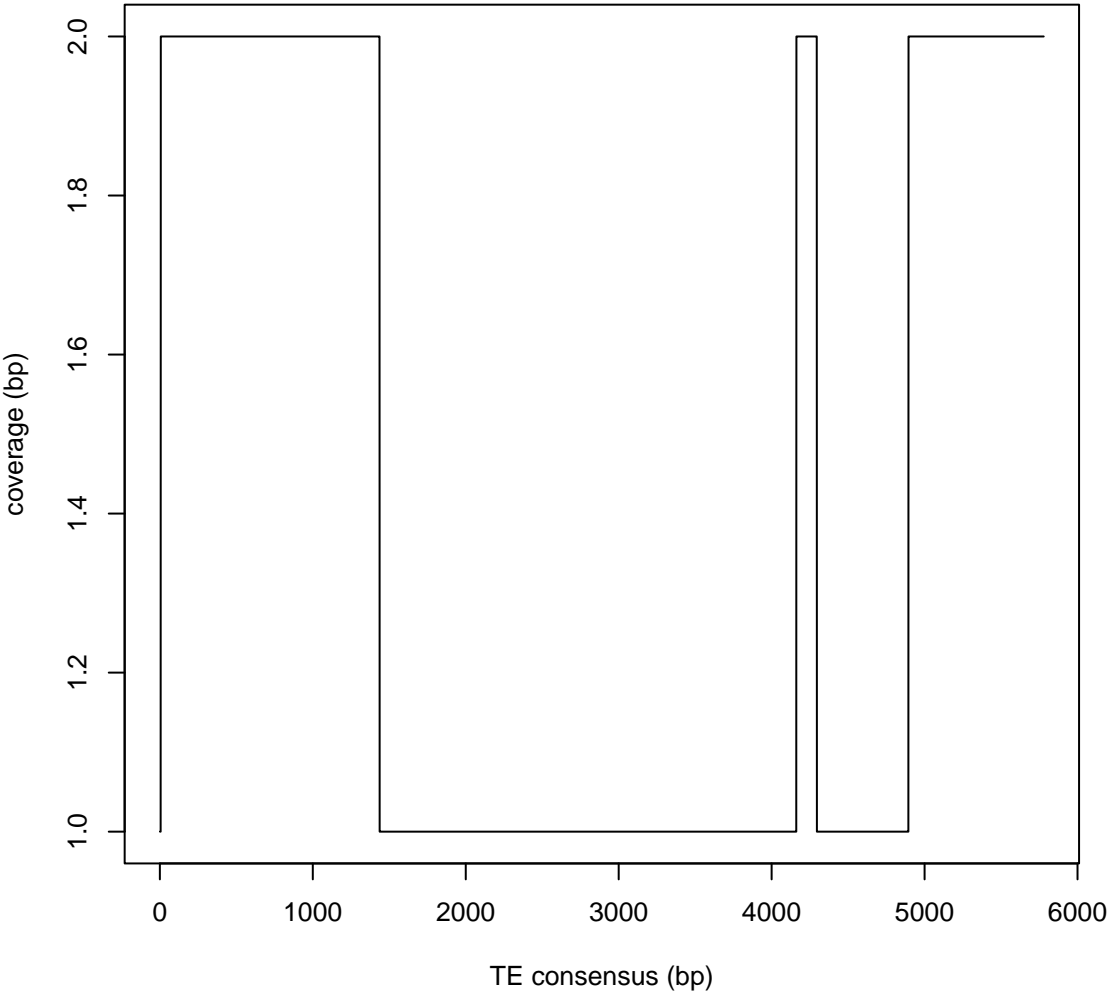


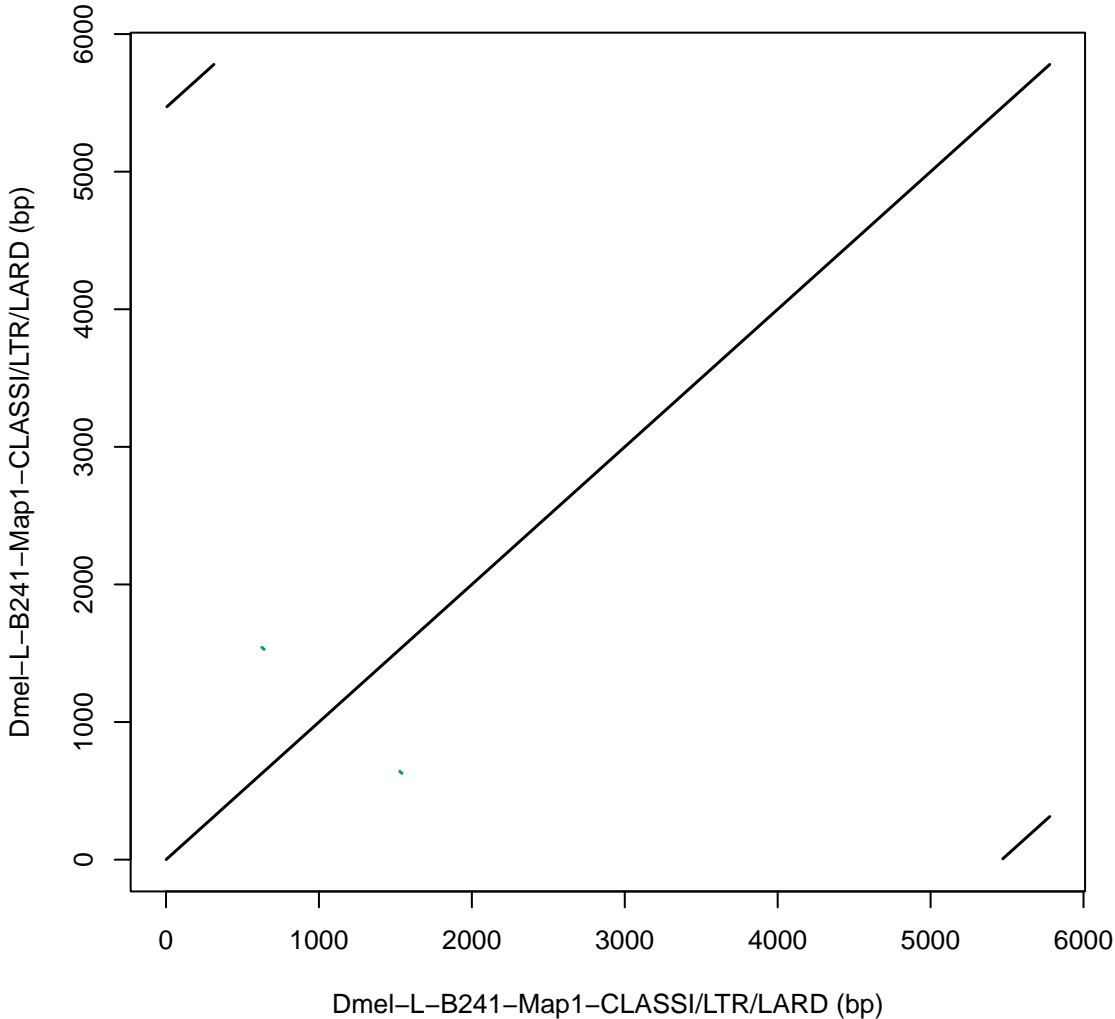
TE: Dmel-L-B241-Map1-CLASSI/LTR/LARD
consensus size: 5779bp; fragments: 4; full length: 1 (>=5201.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

