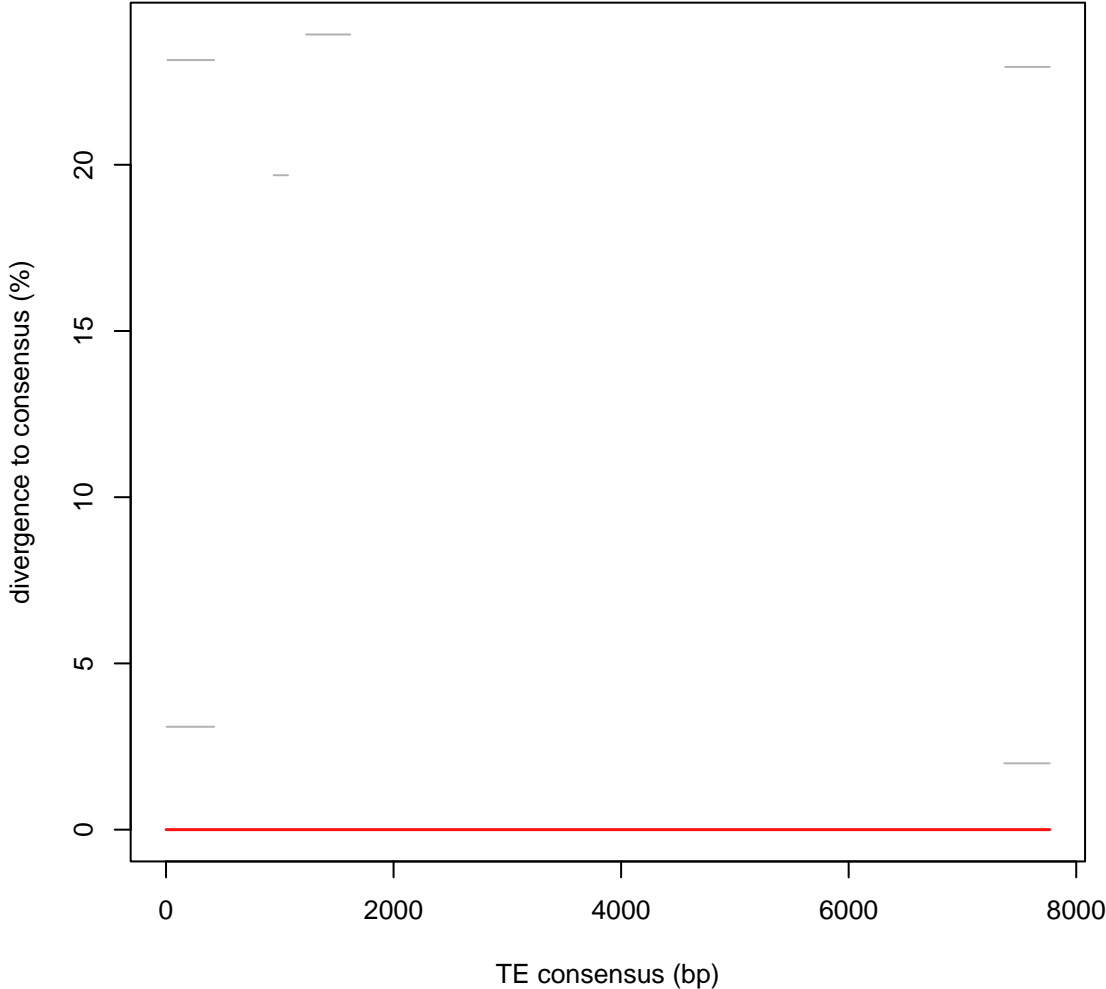
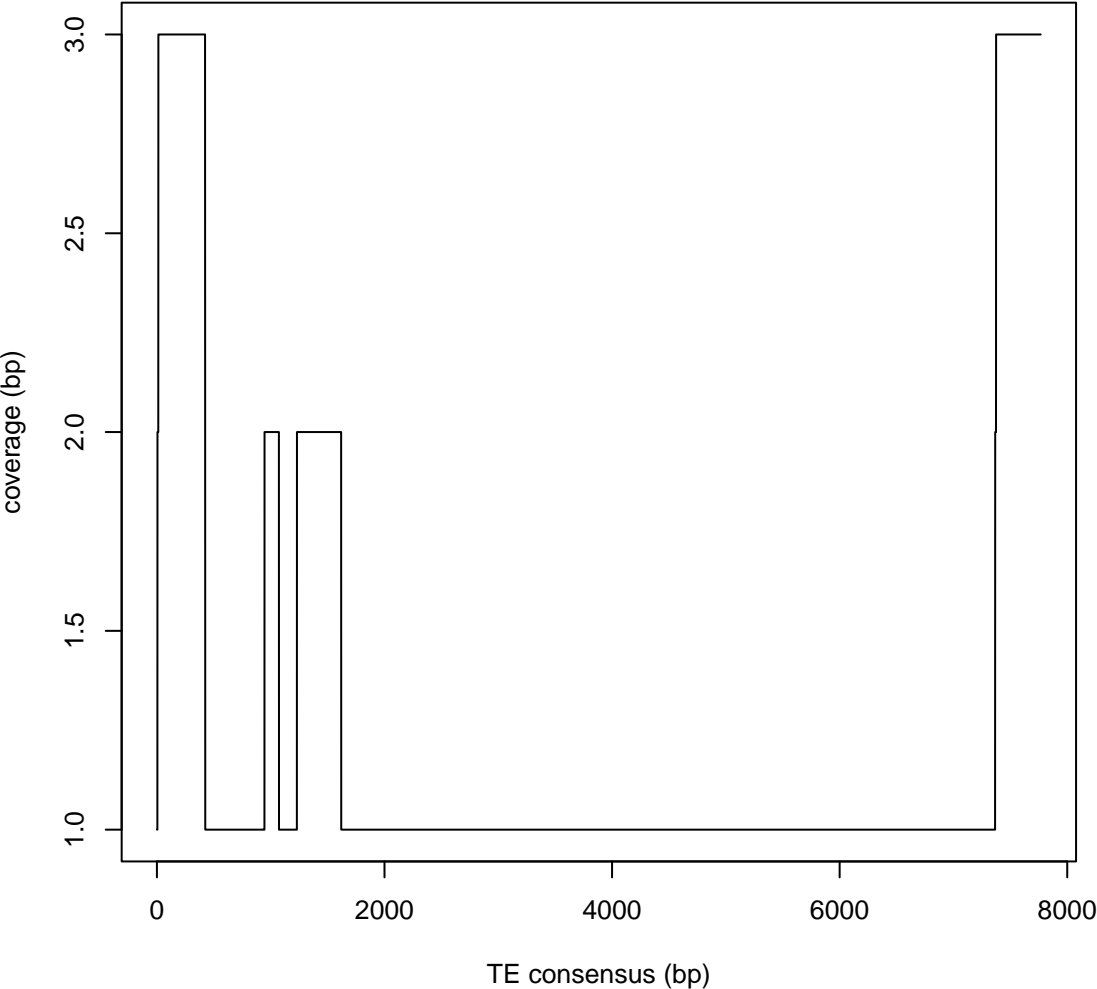


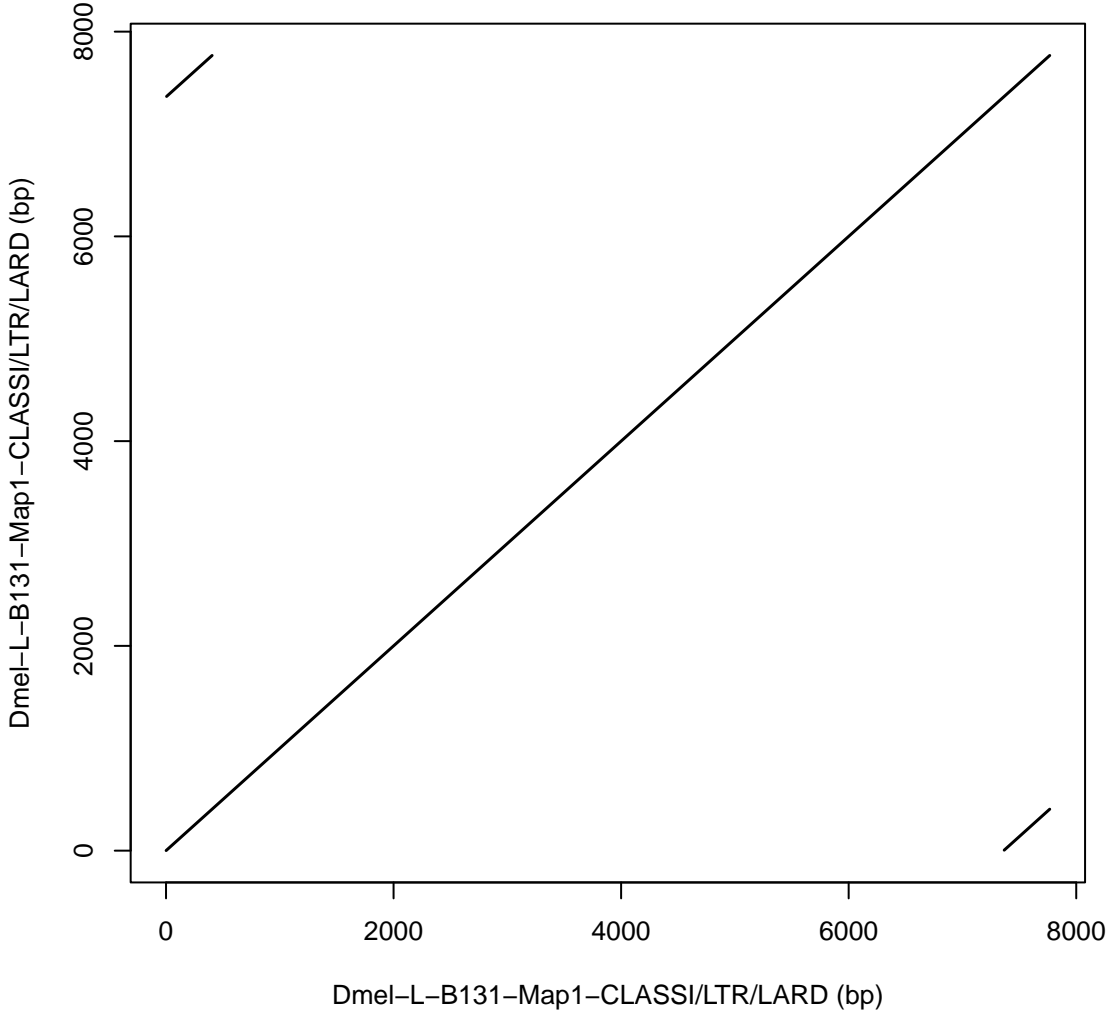
TE: Dmel-L-B131-Map1-CLASSI/LTR/LARD
consensus size: 7767bp; fragments: 7; full length: 1 (>=6990.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

