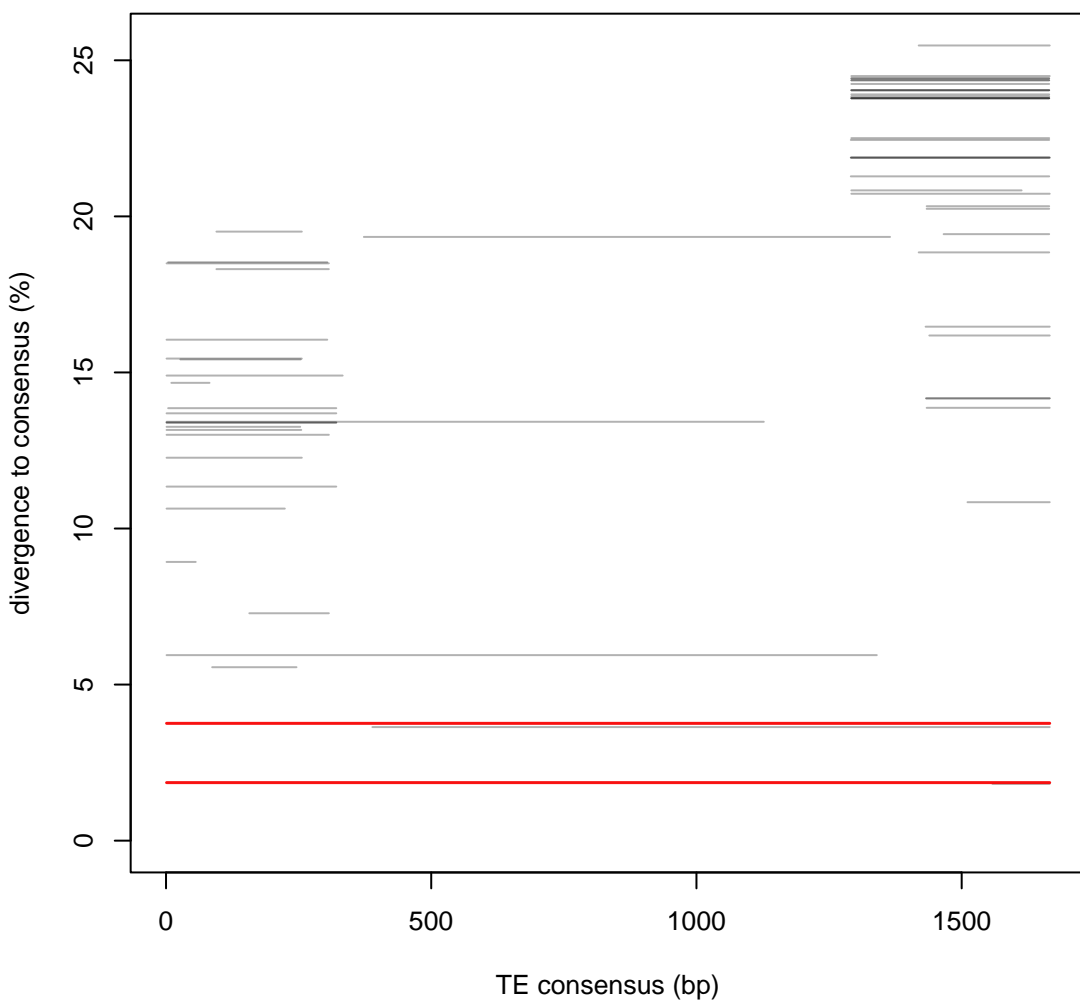
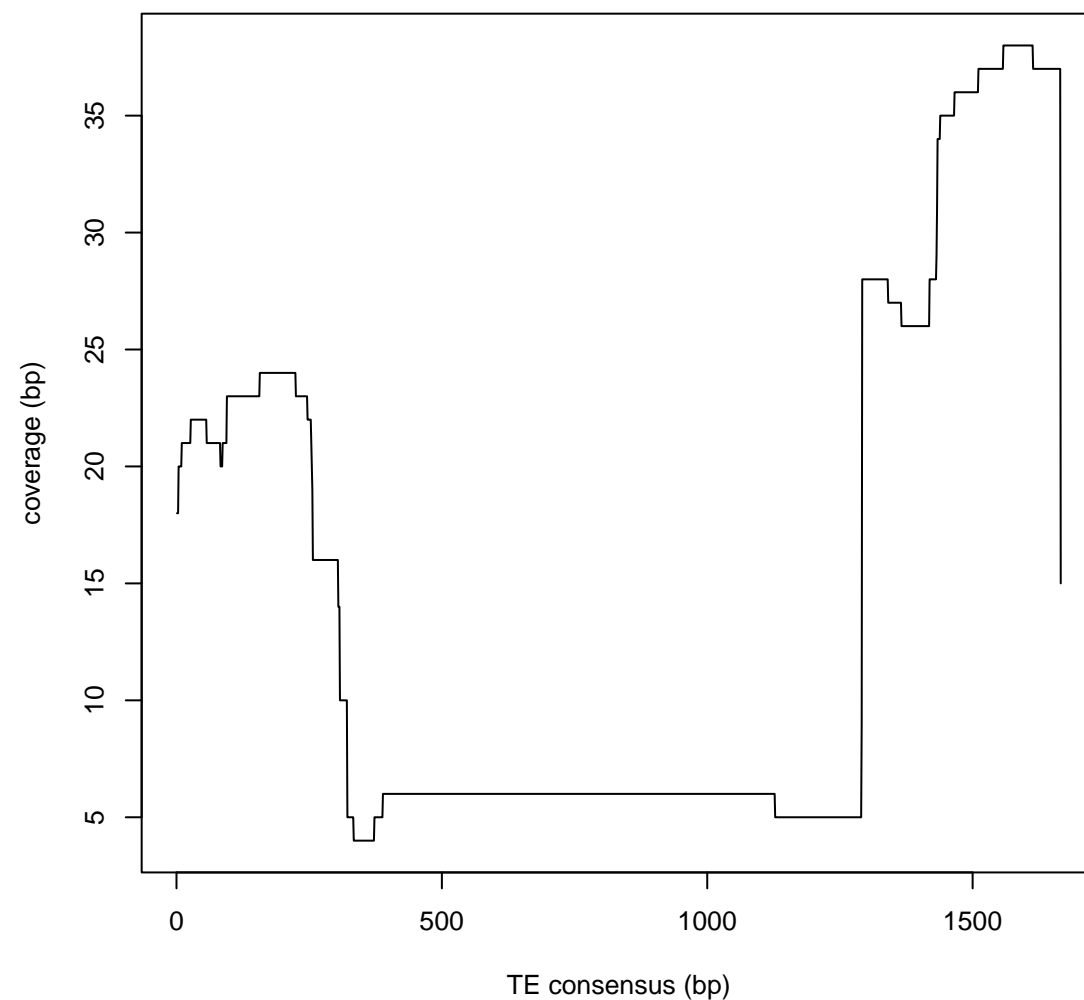


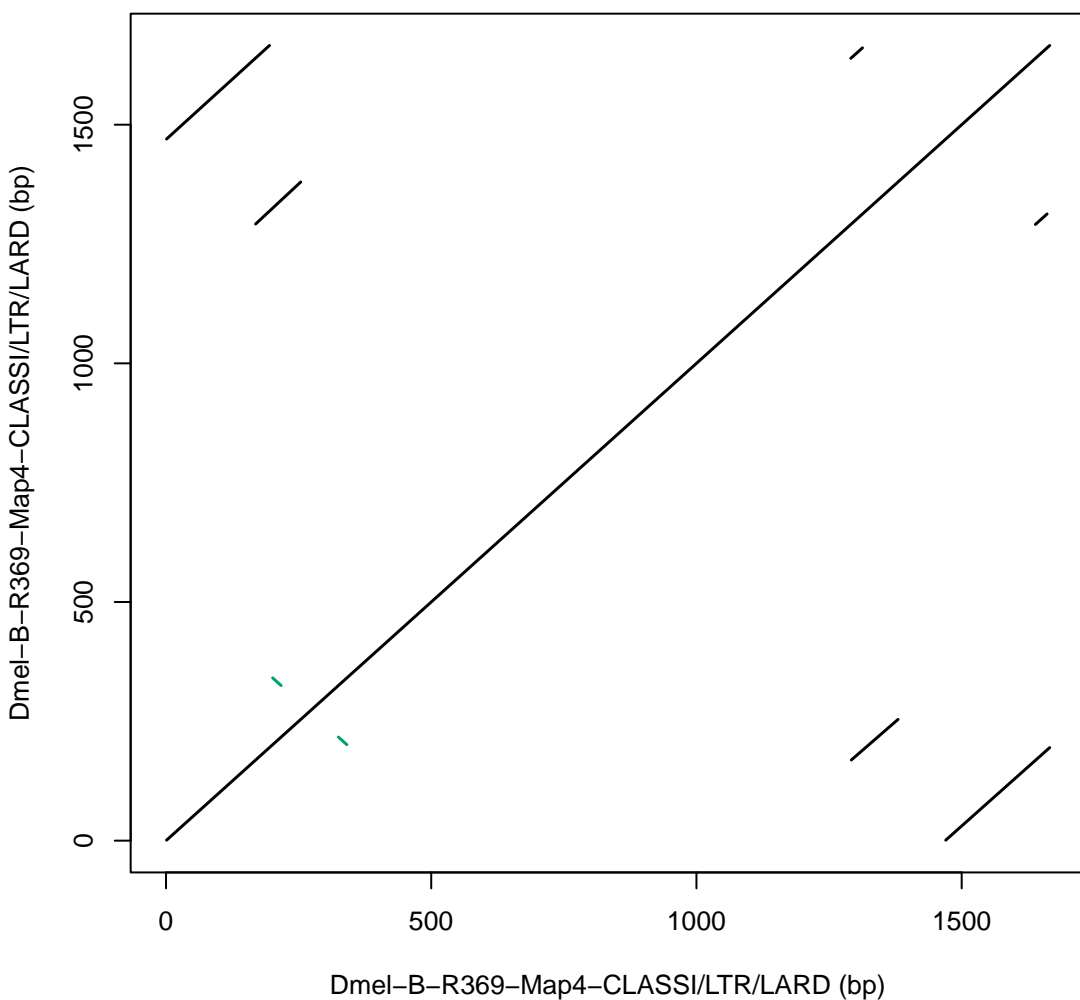
TE: Dmel-B-R369-Map4-CLASSI/LTR/LARD  
consensus size: 1666bp; fragments: 63; full length: 2 (>=1499.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

