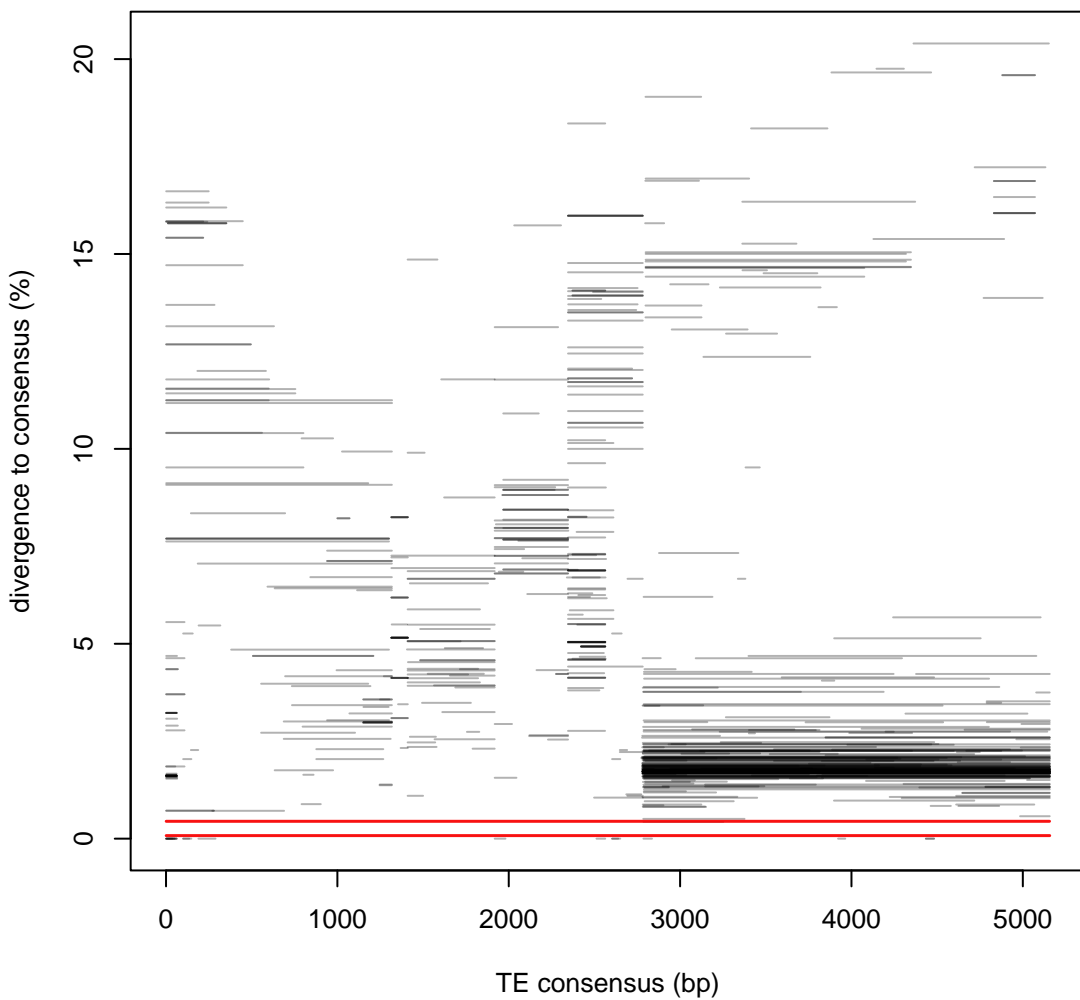
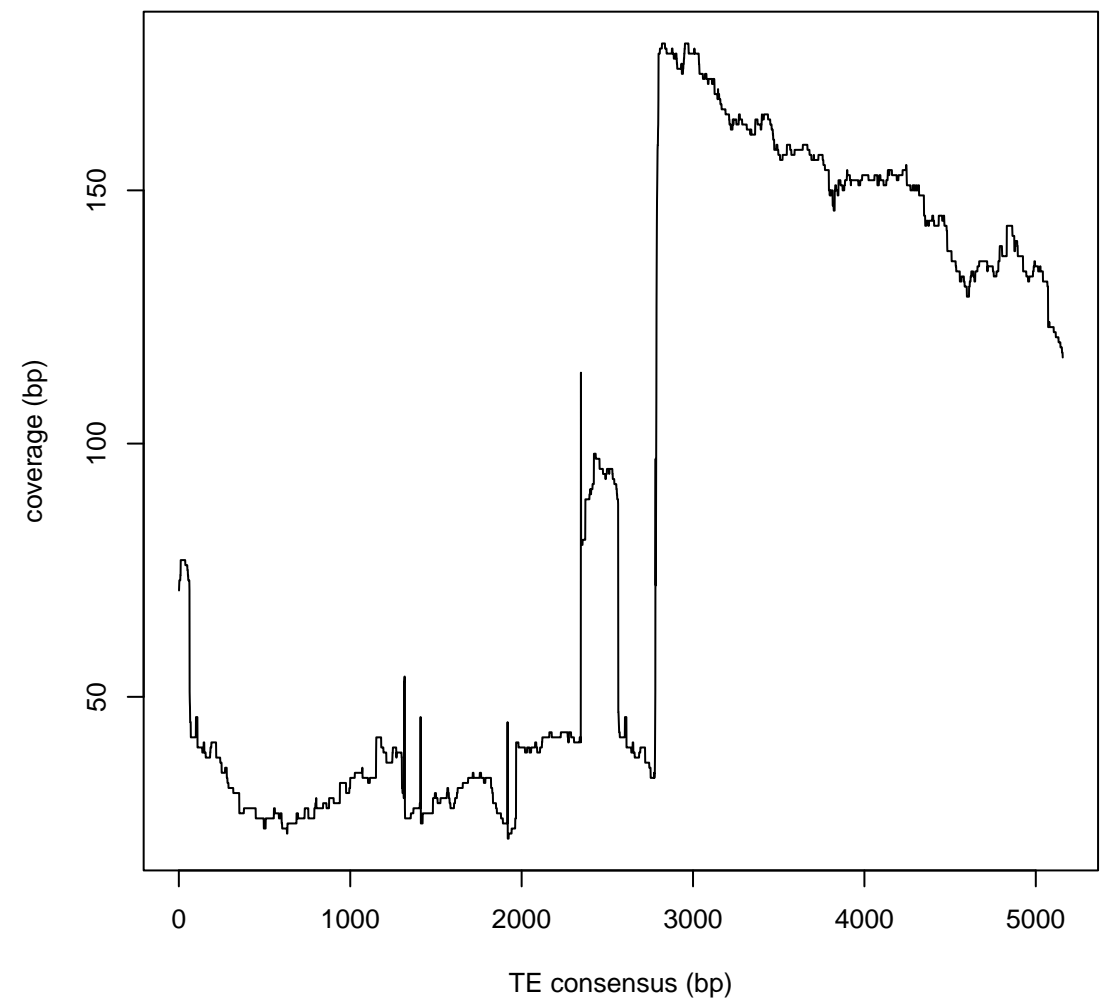


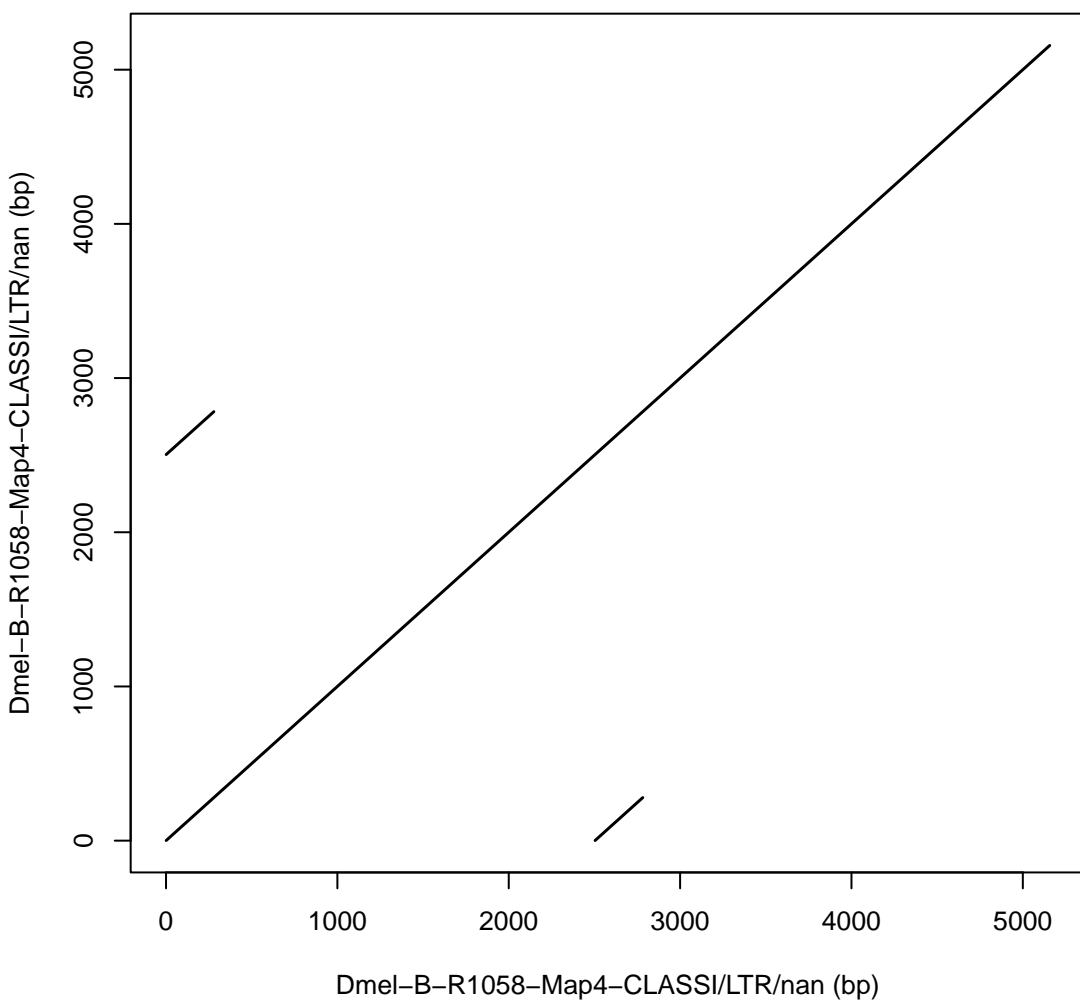
TE: Dmel-B-R1058-Map4-CLASSI/LTR/nan  
consensus size: 5157bp; fragments: 639; full length: 2 (>=4641.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

