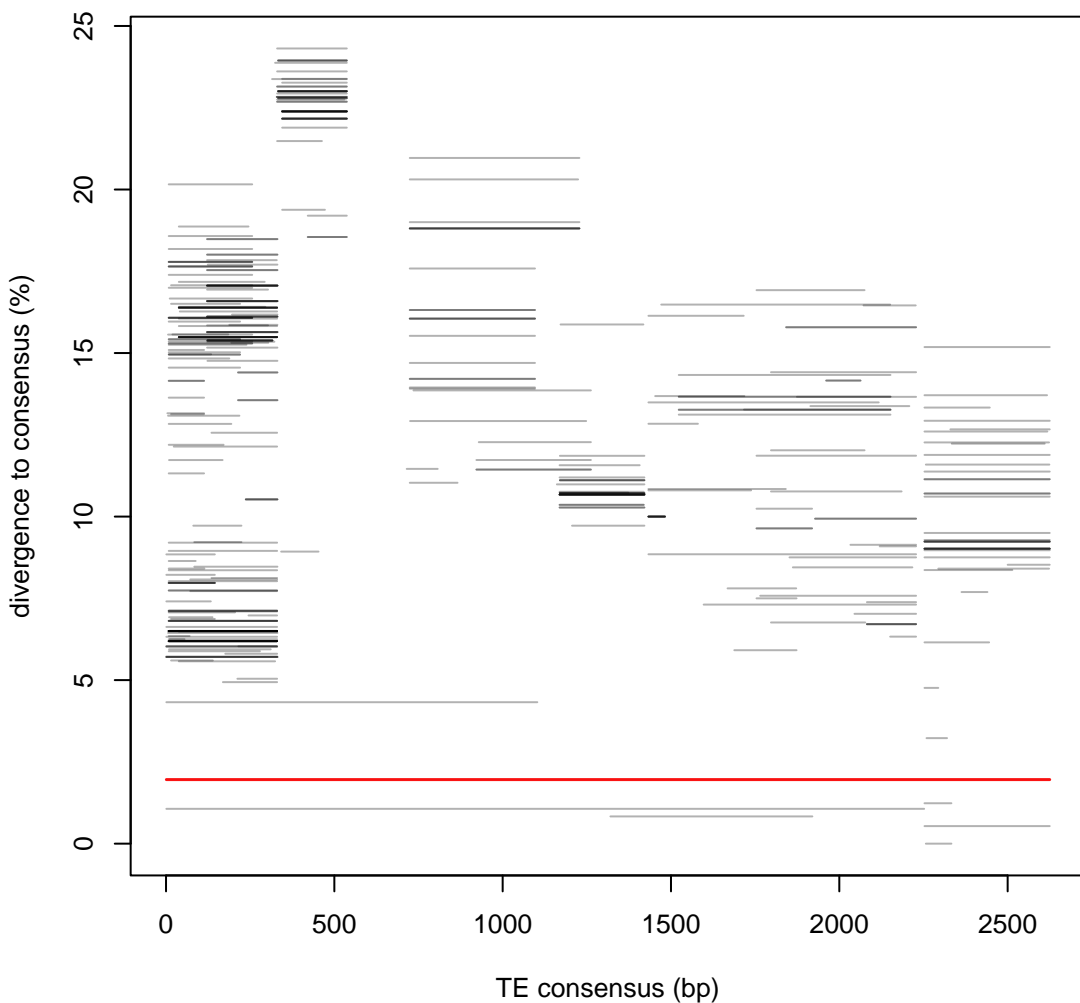
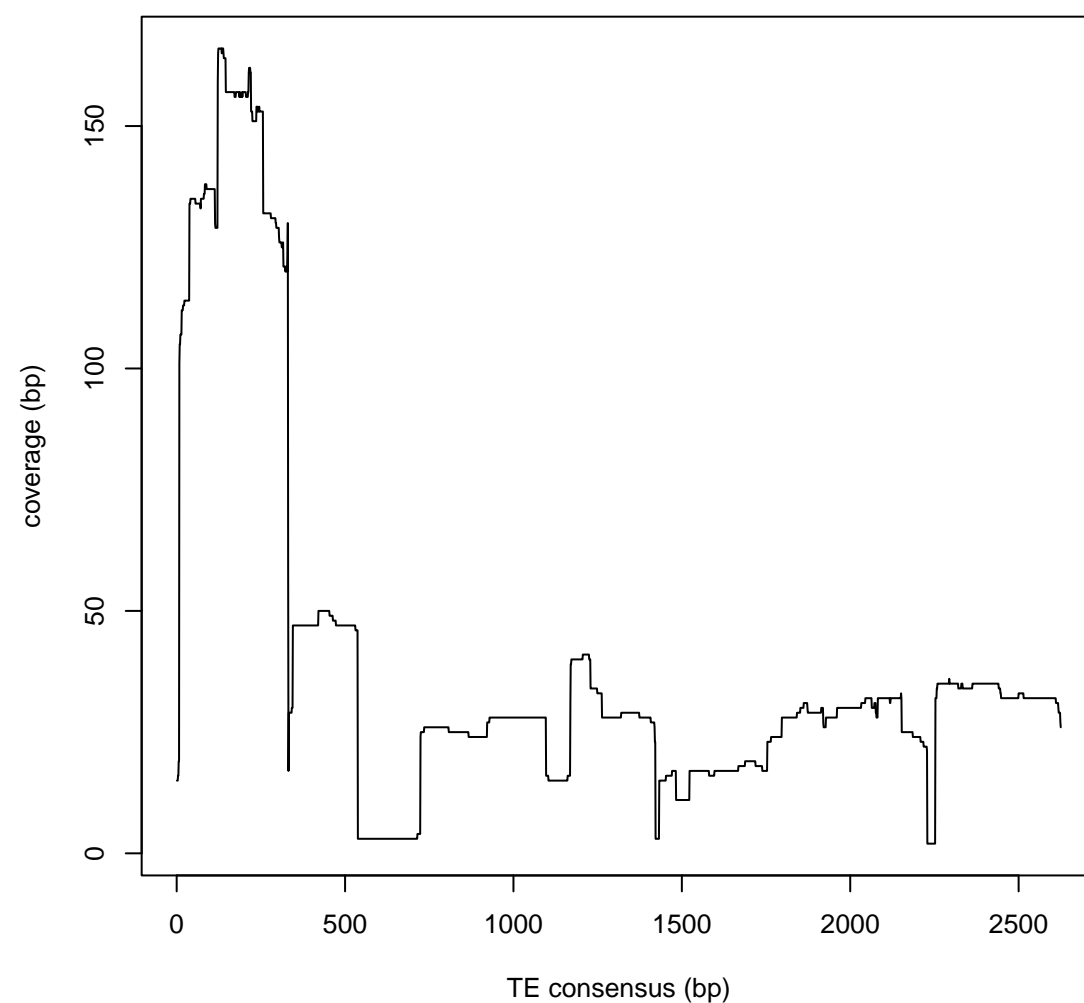


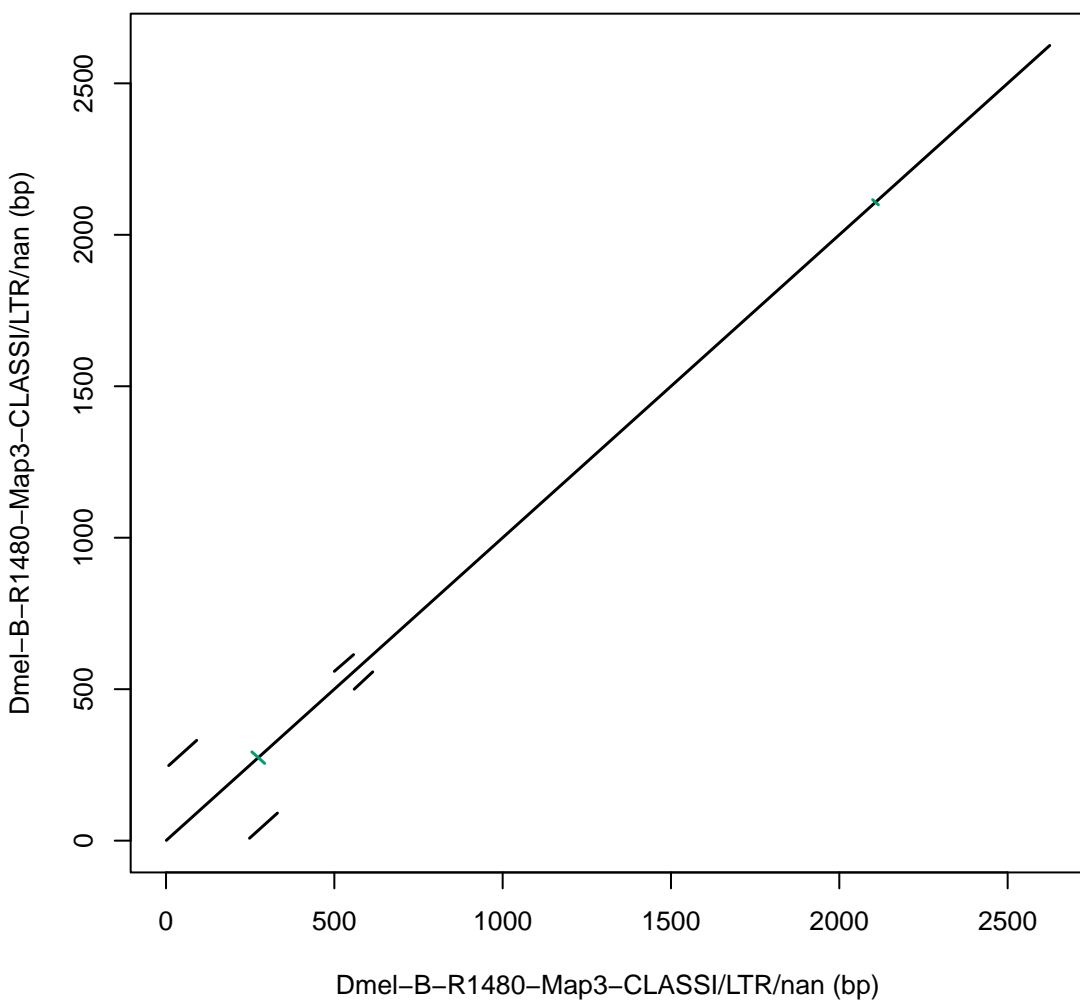
TE: Dmel-B-R1480-Map3-CLASSI/LTR/nan  
consensus size: 2625bp; fragments: 388; full length: 1 (>=2362.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

