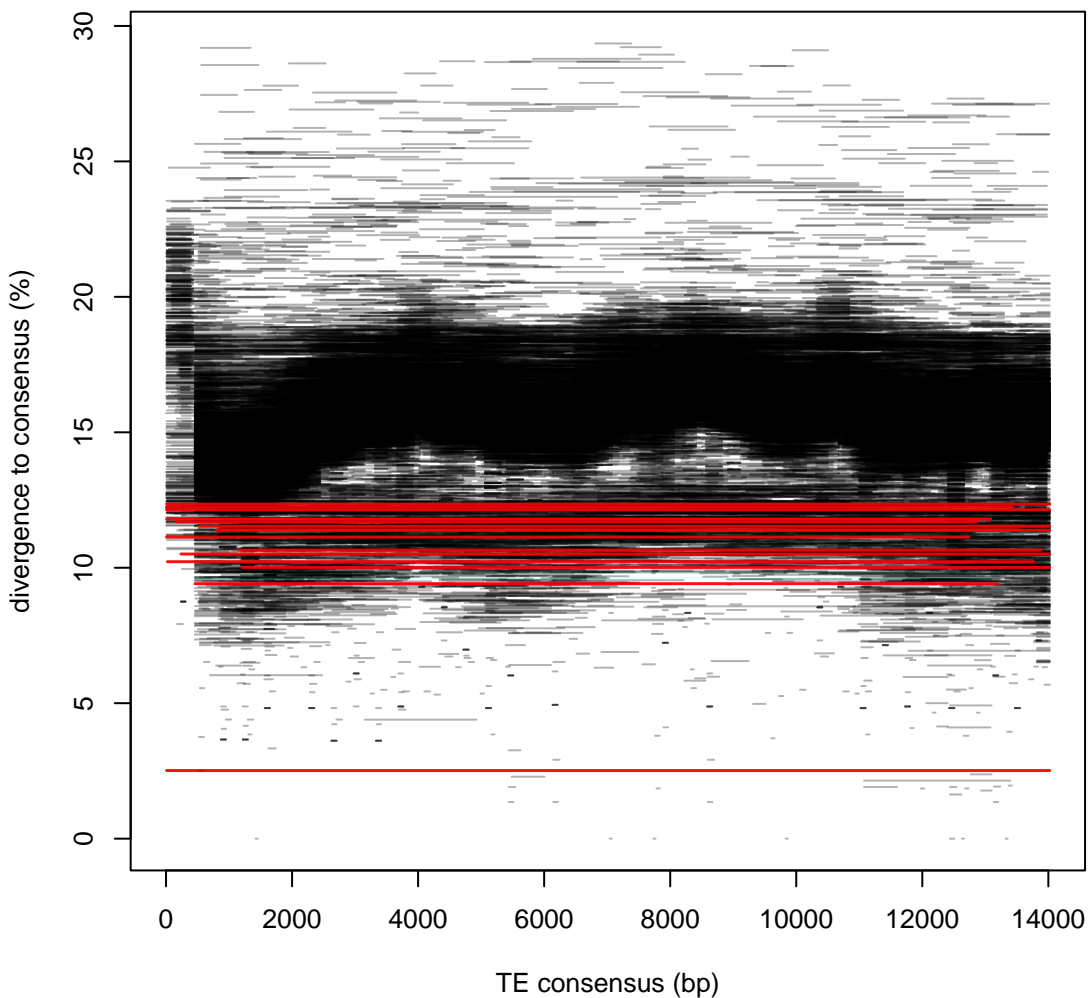
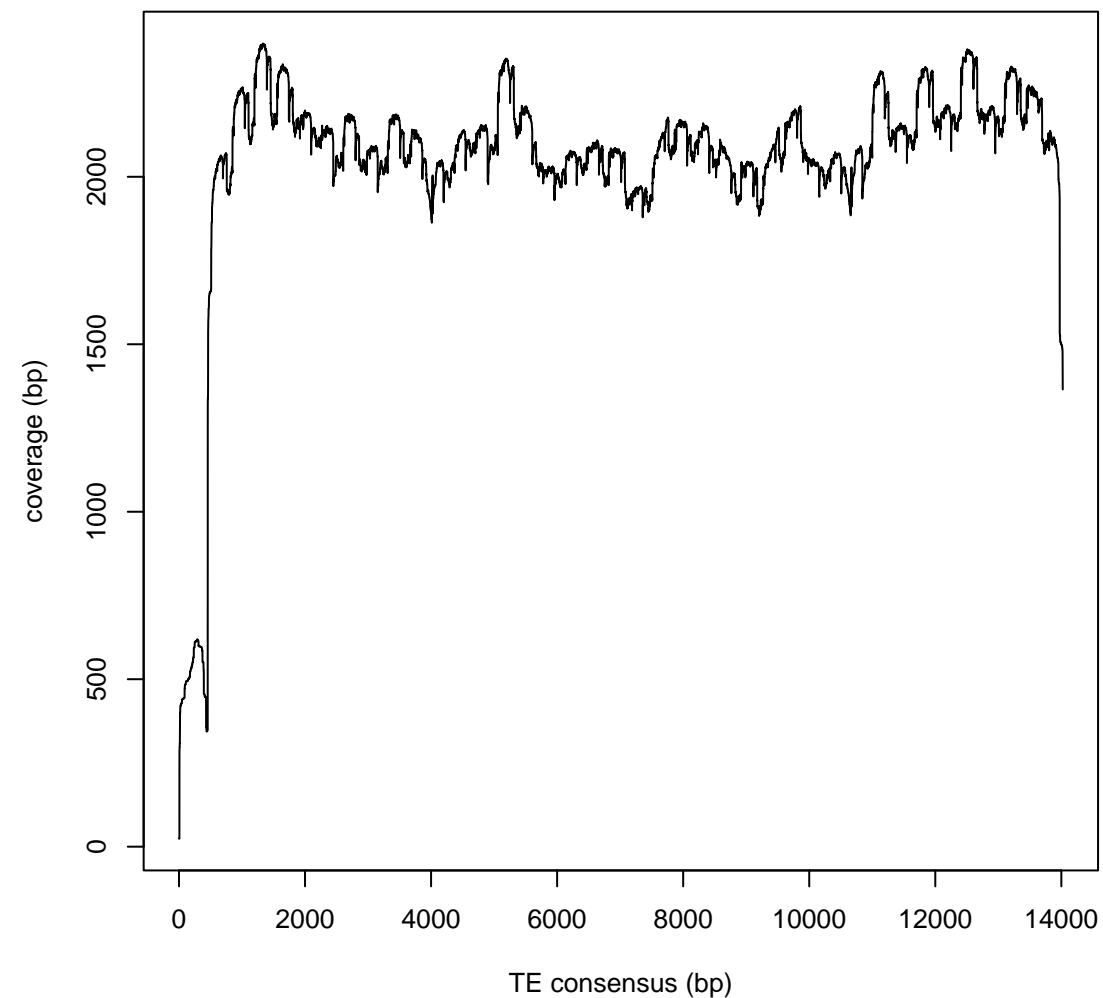


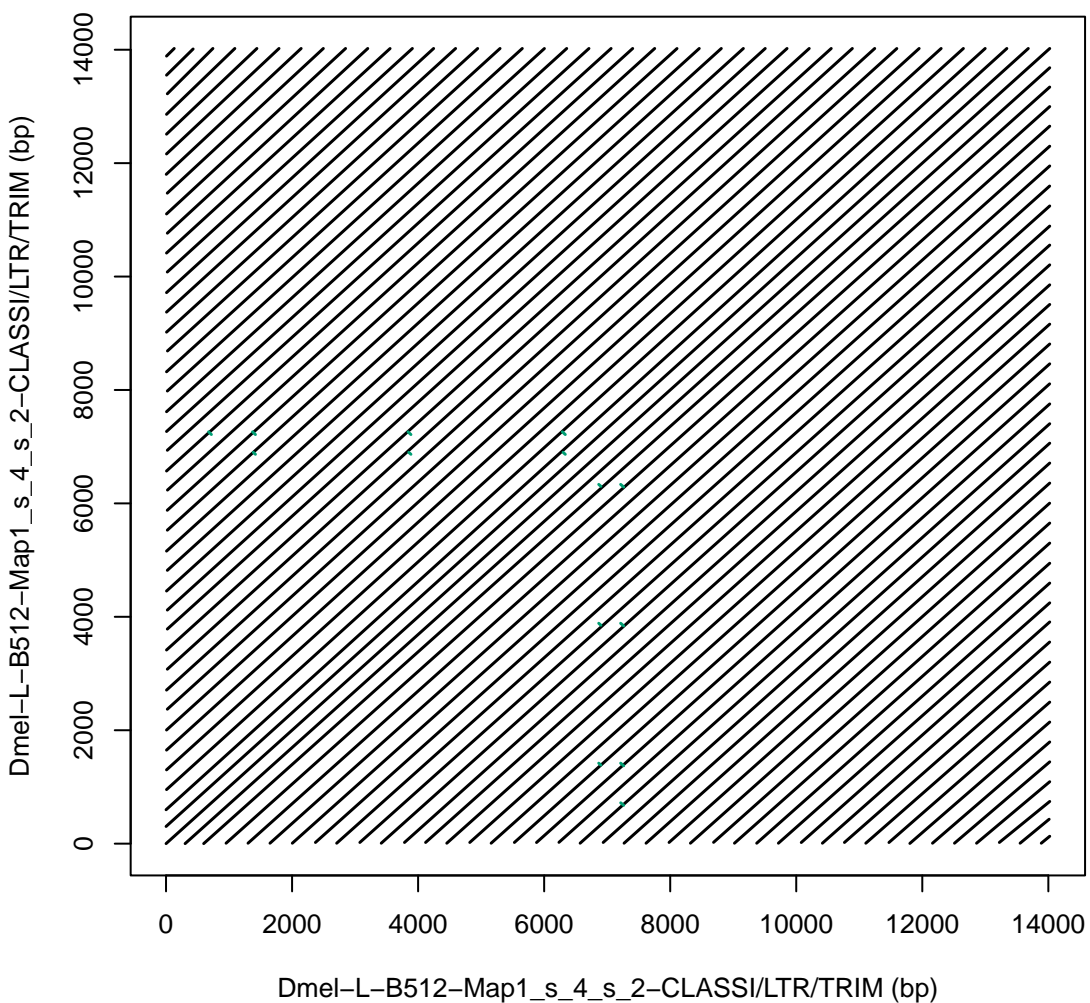
TE: Dmel-L-B512-Map1_s_4_s_2-CLASSI/LTR/TRIM
consensus size: 14022bp; fragments: 27063; full length: 15 (>=12619.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

