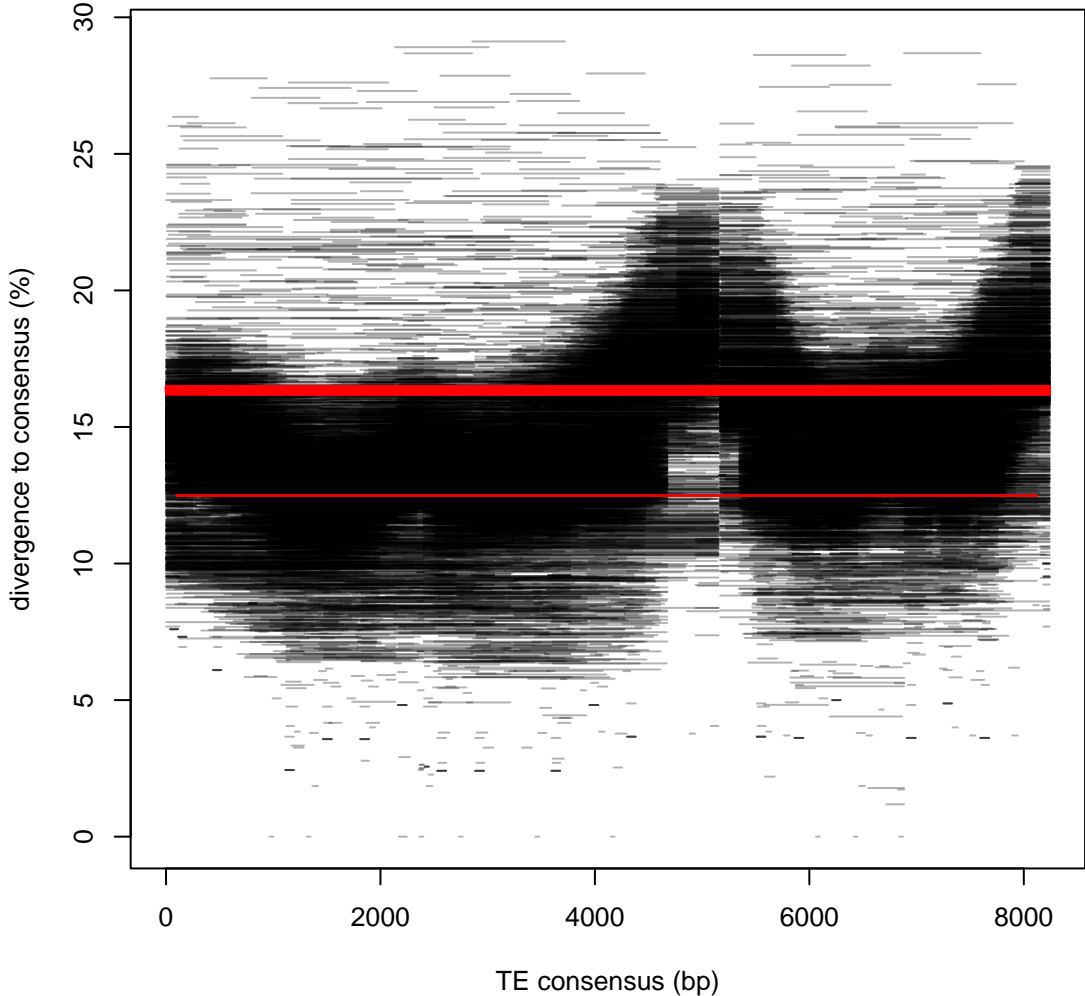
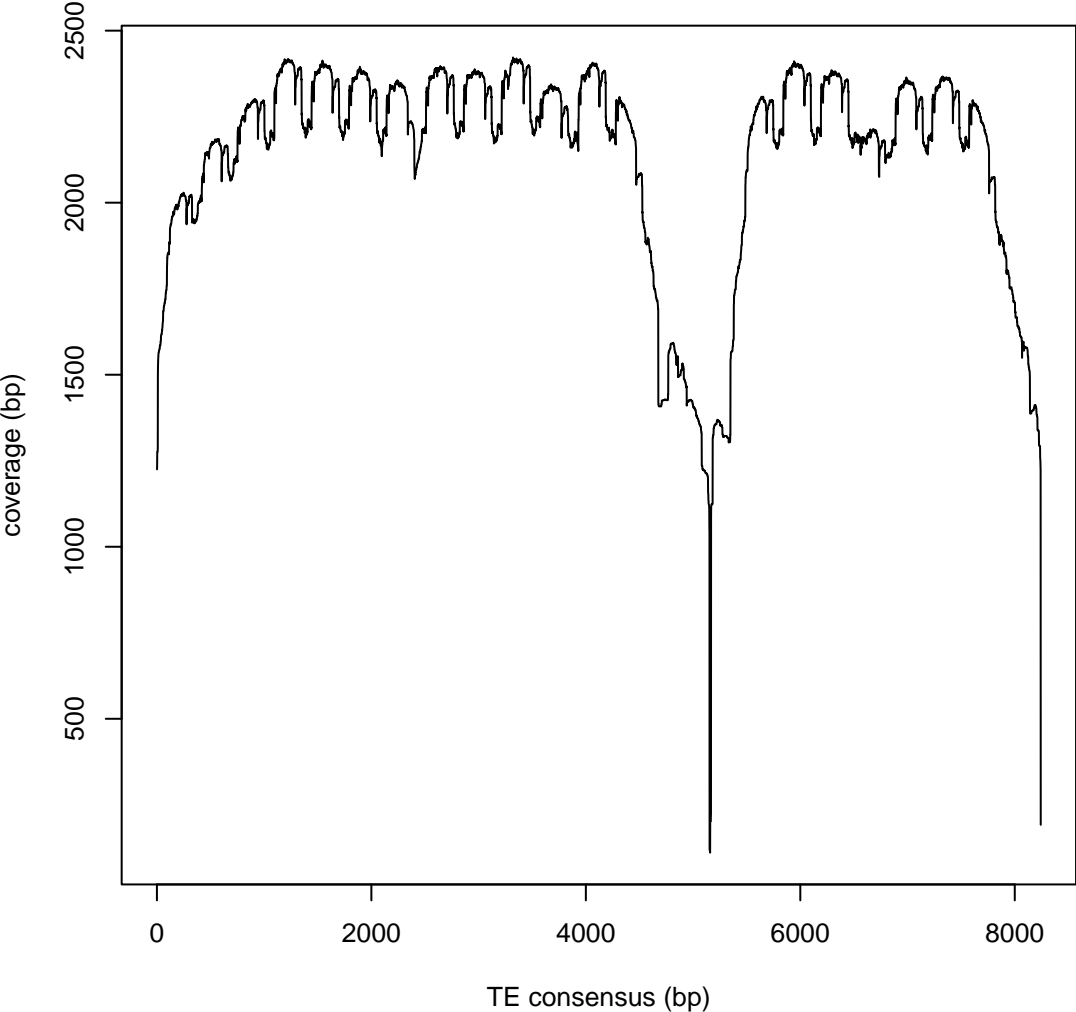


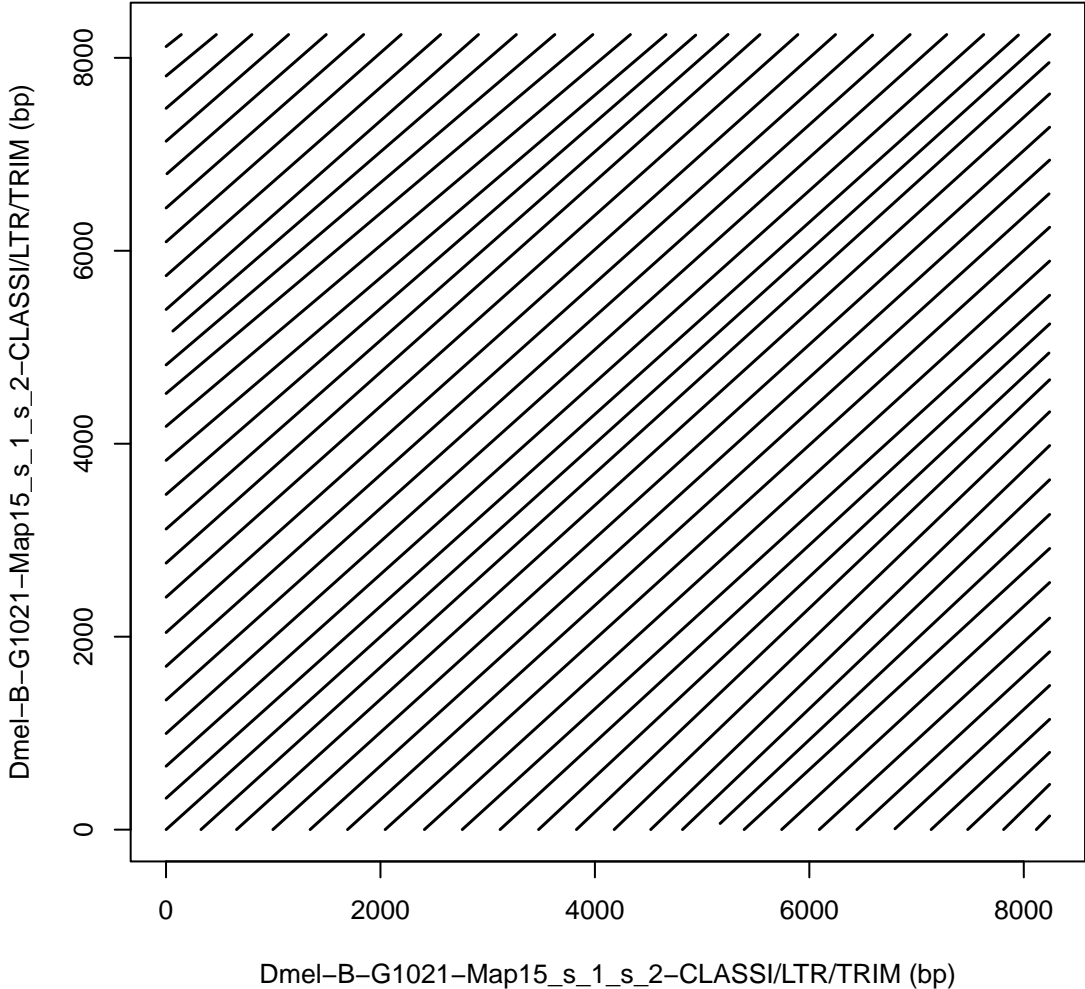
TE: Dmel-B-G1021-Map15_s_1_s_2-CLASSI/LTR/TRIM
consensus size: 8242bp; fragments: 19286; full length: 13 (>=7417.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

