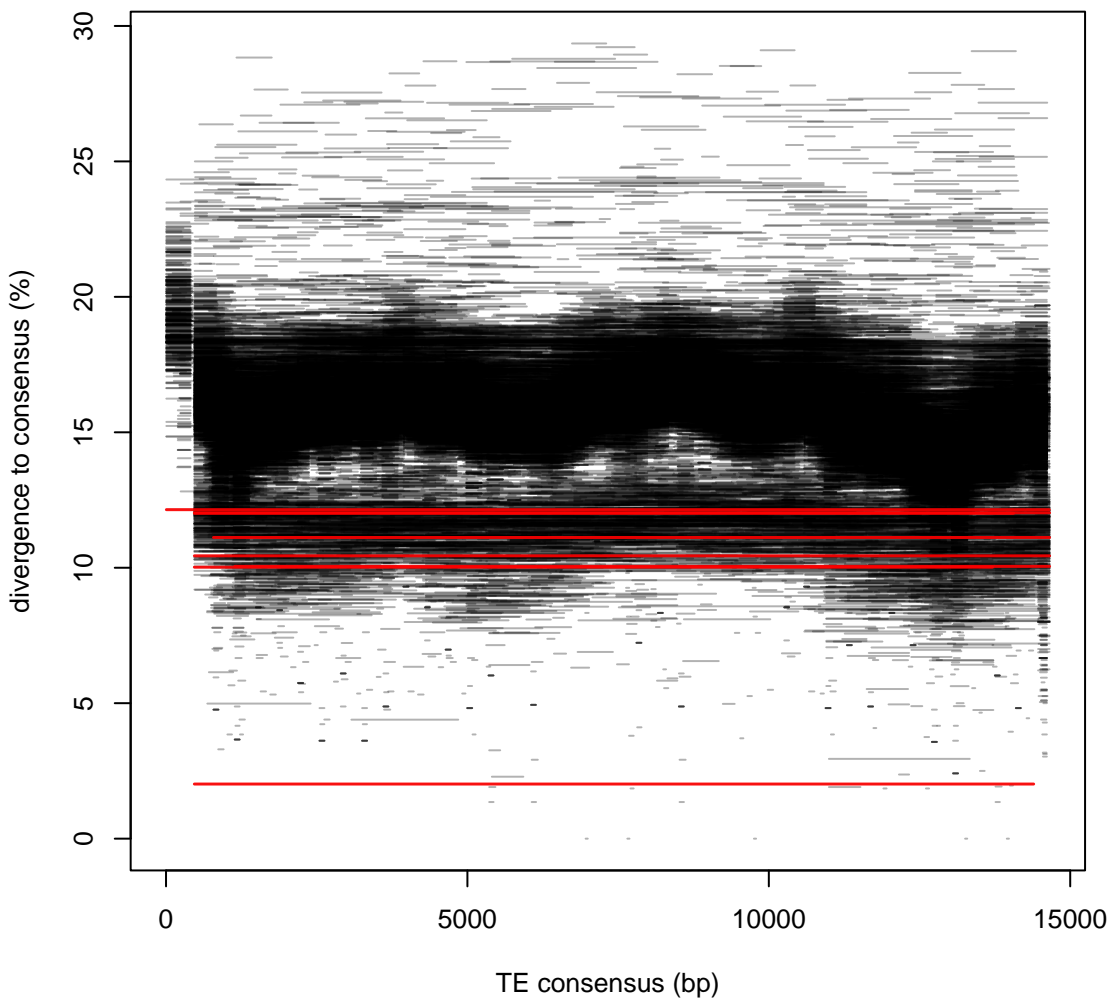
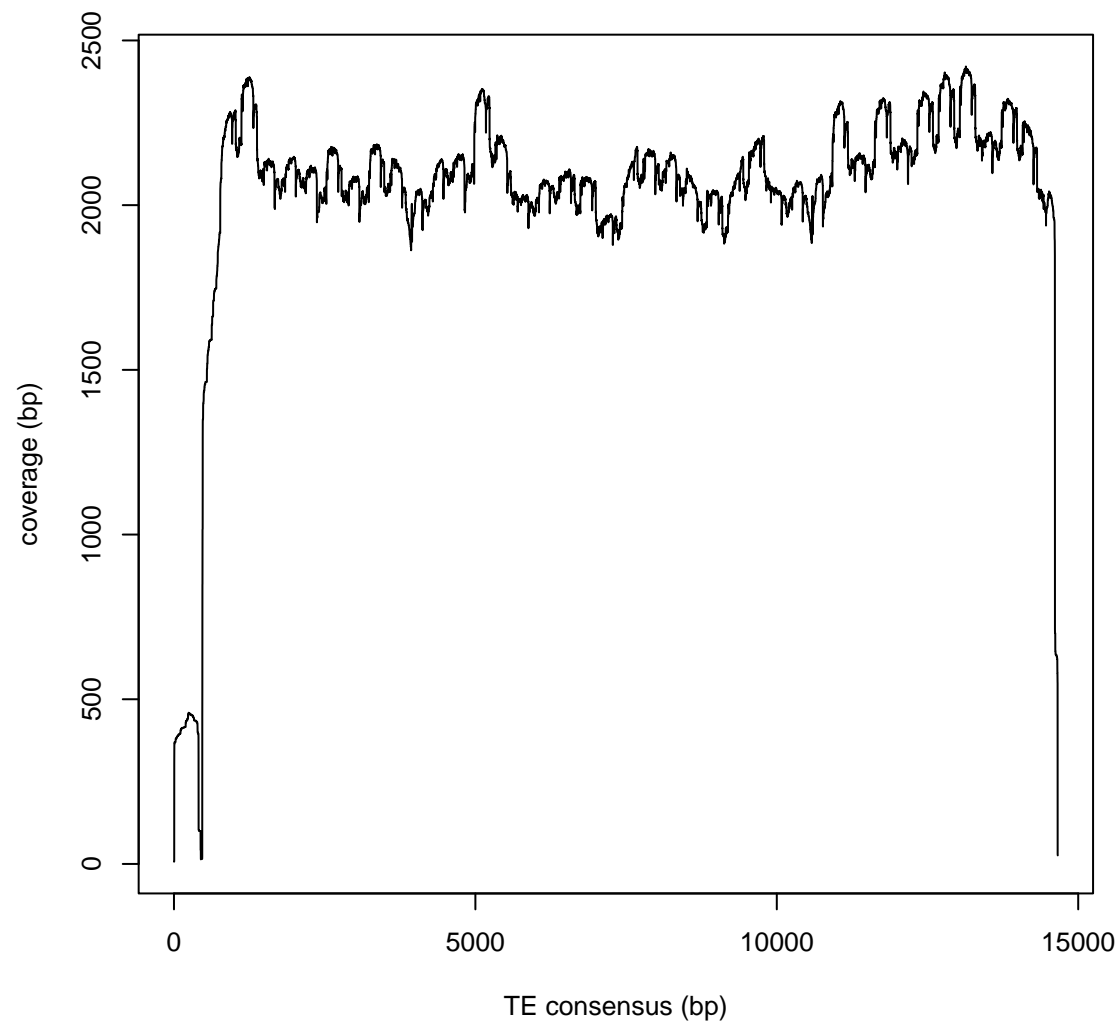


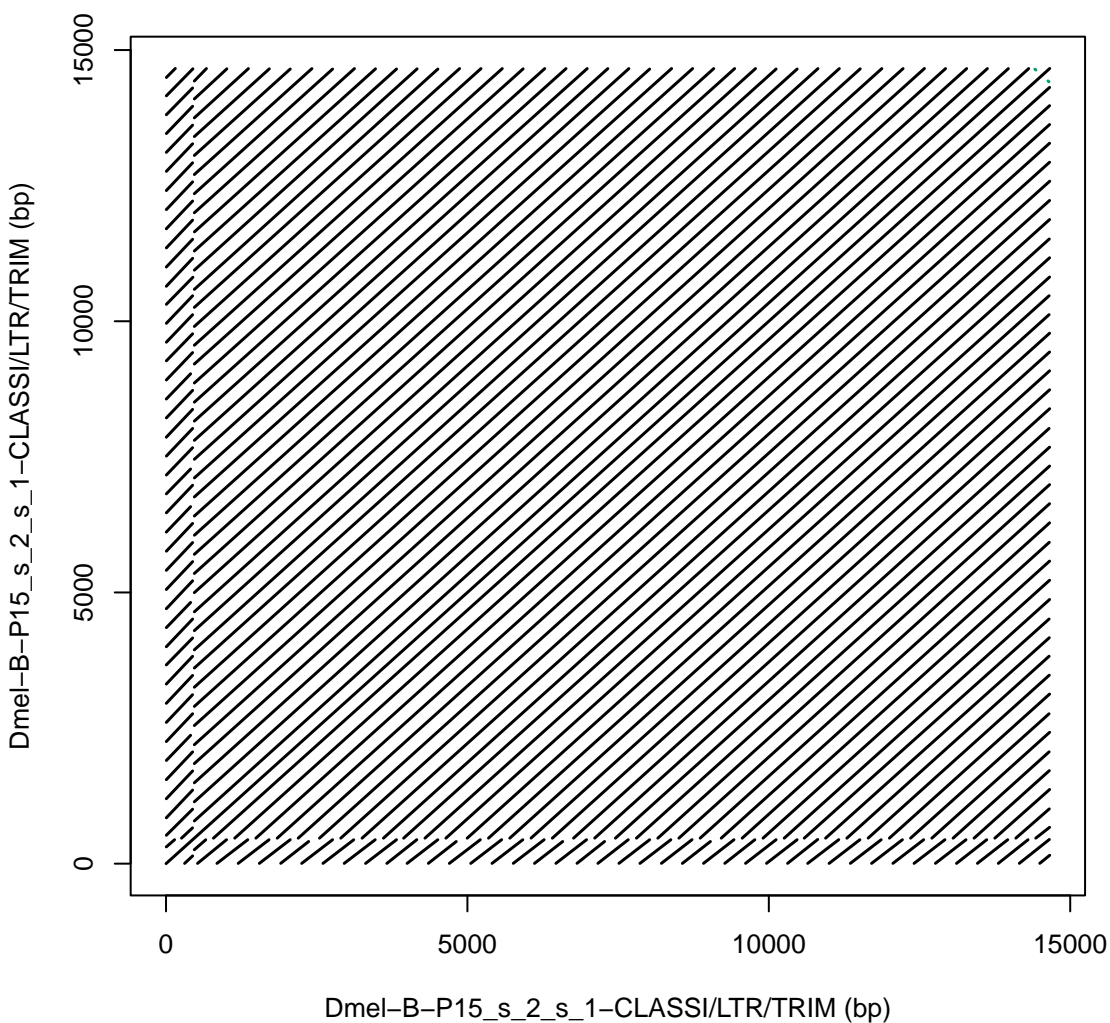
TE: Dmel-B-P15_s_2_s_1-CLASSI/LTR/TRIM
consensus size: 14660bp; fragments: 28014; full length: 8 (>=13194bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

