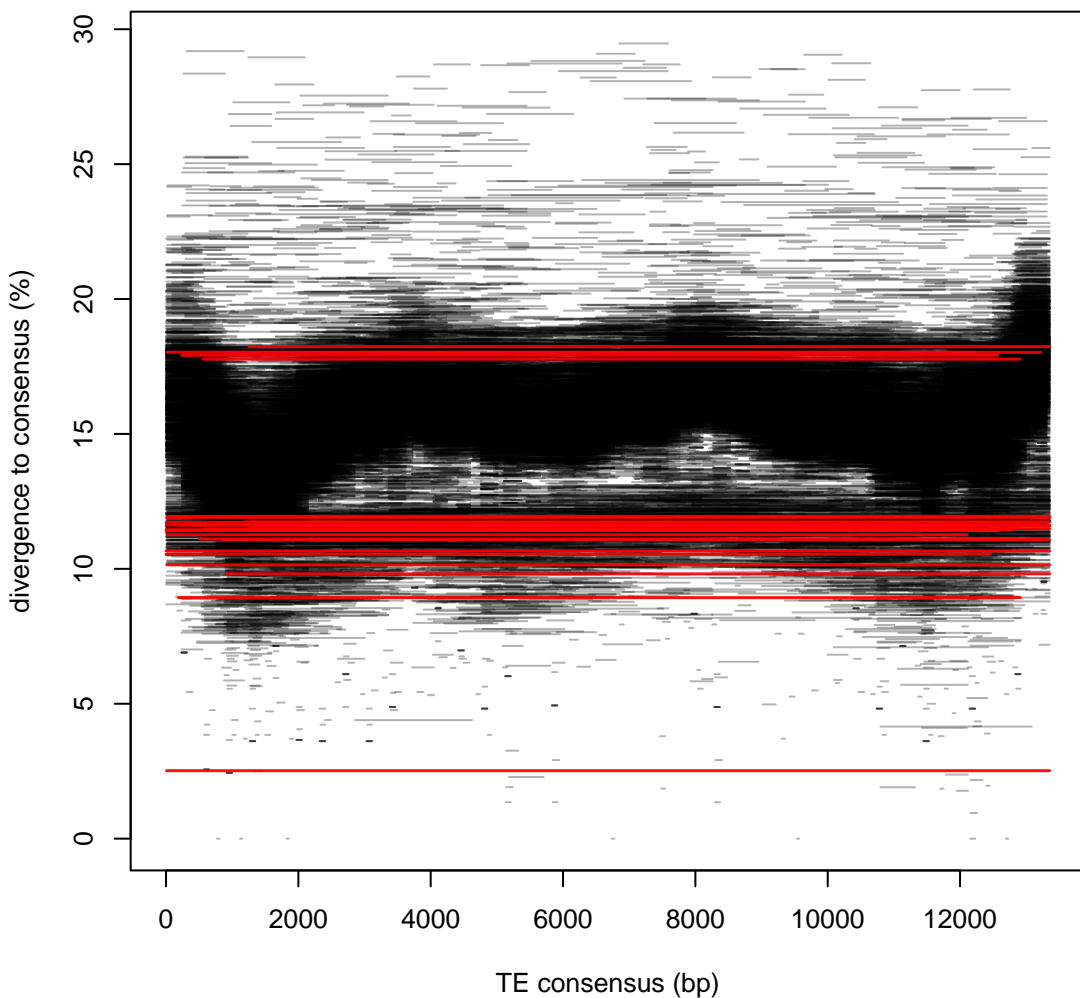
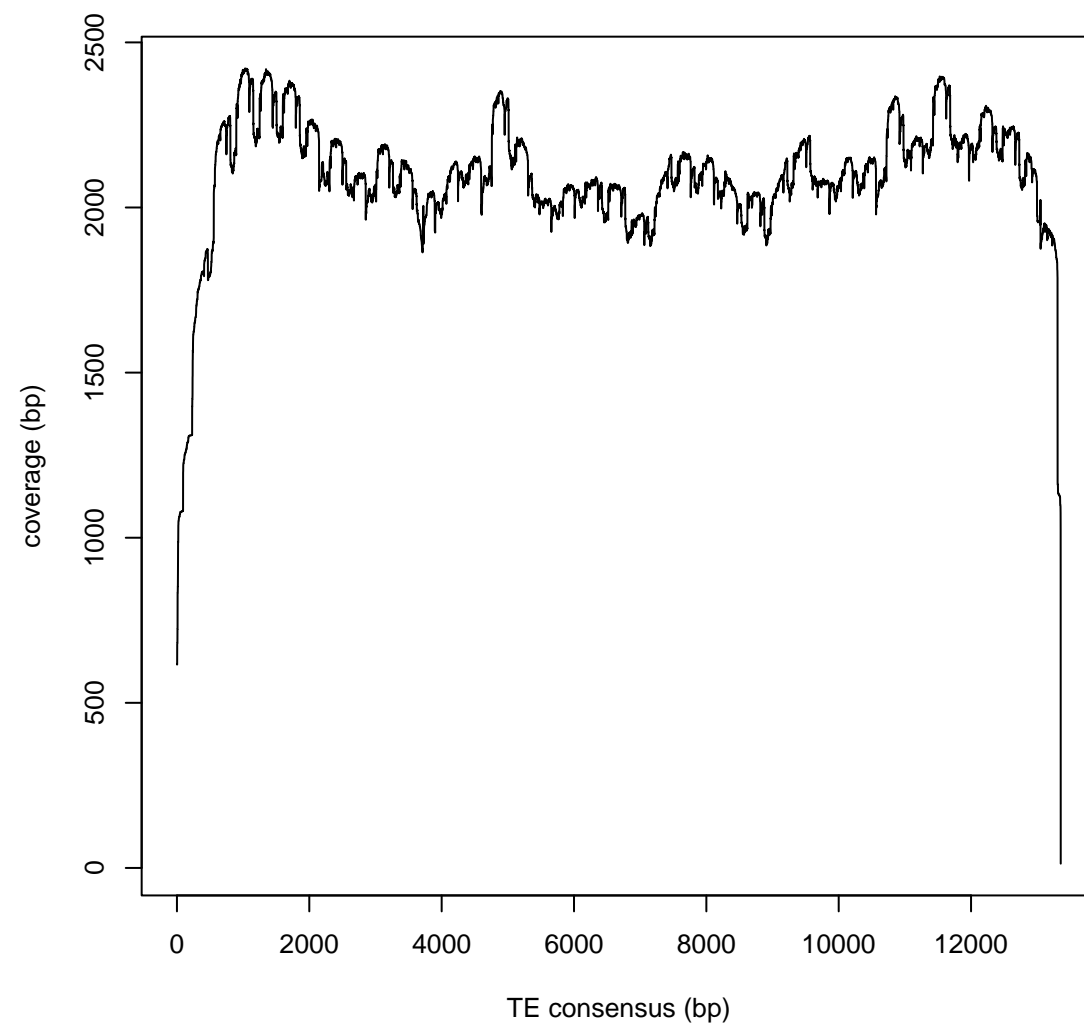


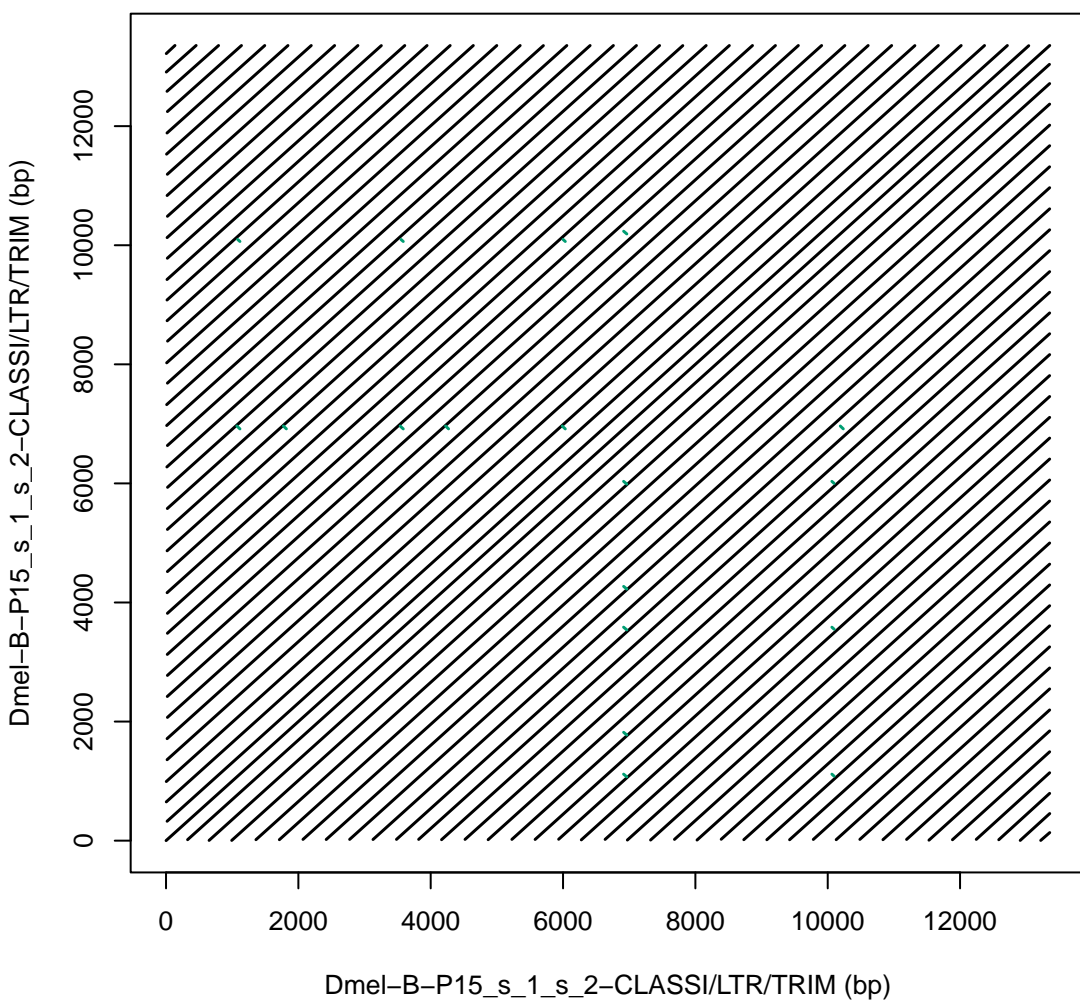
TE: Dmel-B-P15_s_1_s_2-CLASSI/LTR/TRIM
consensus size: 13355bp; fragments: 25956; full length: 24 (>=12019.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

