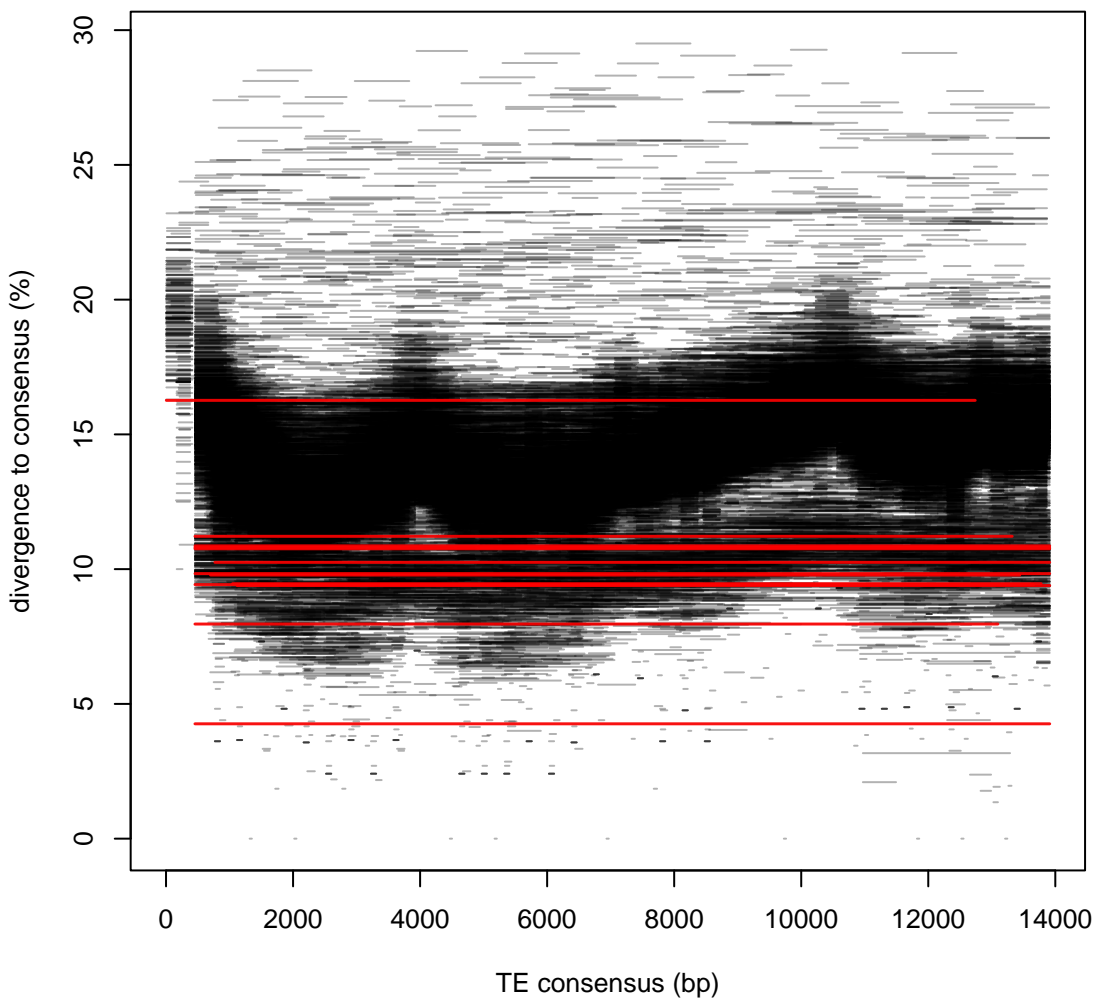
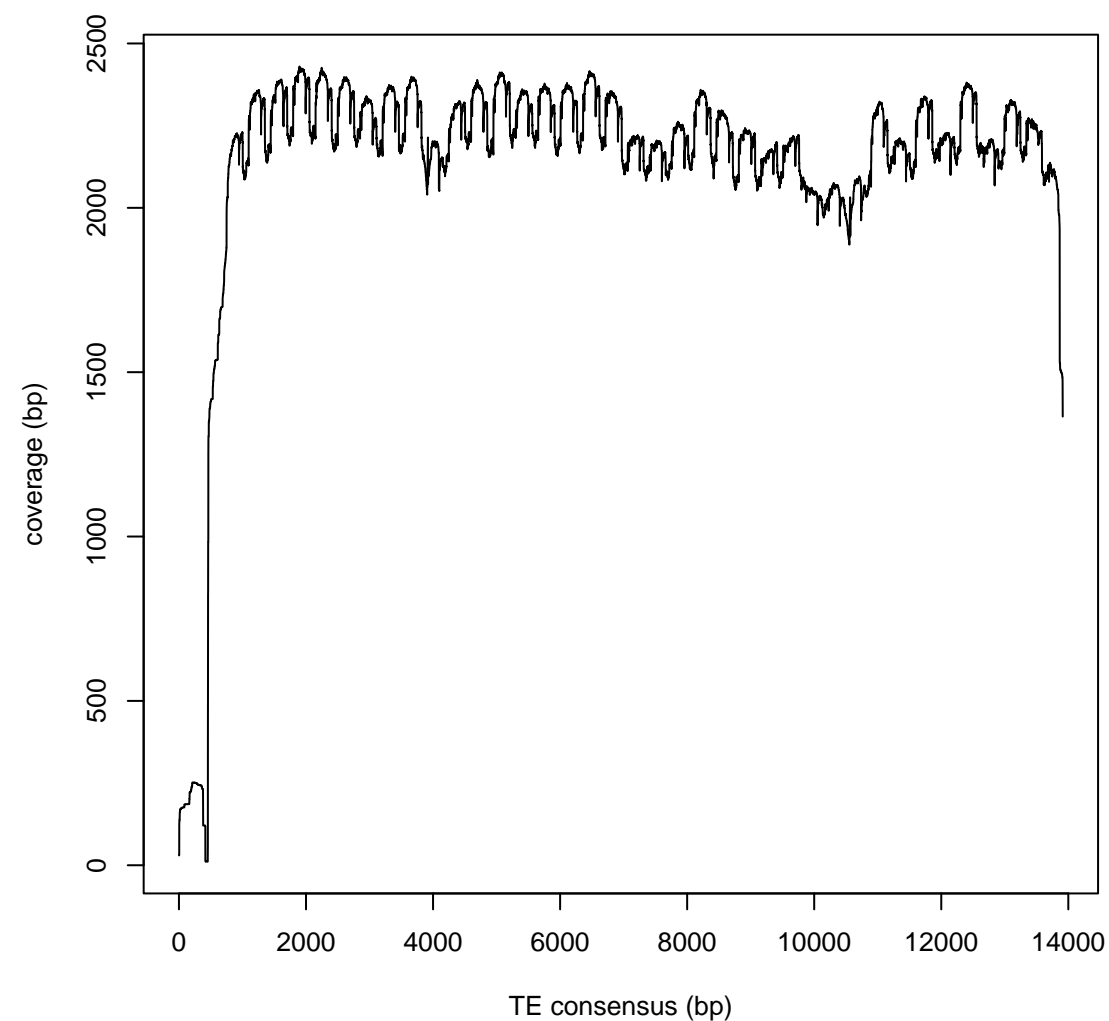


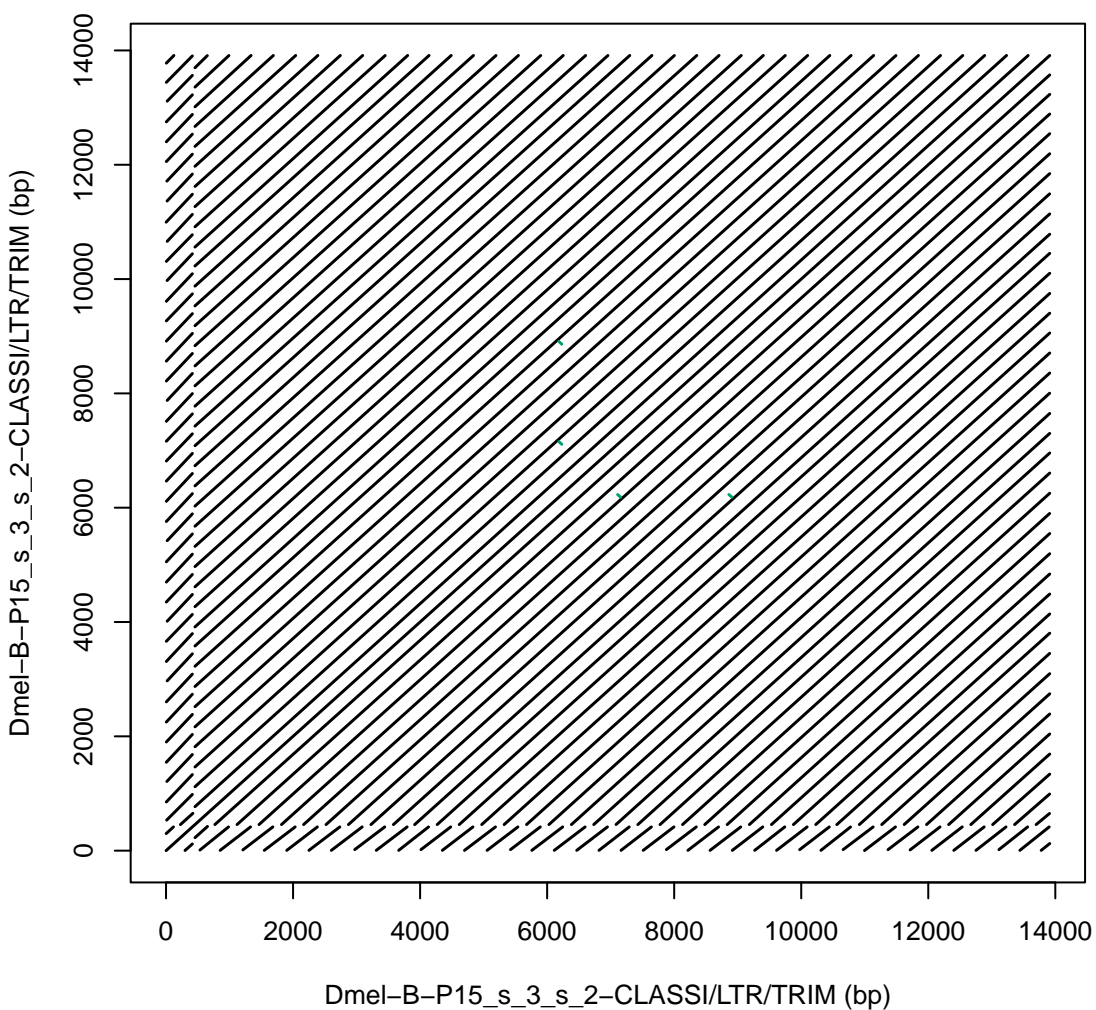
TE: Dmel-B-P15_s_3_s_2-CLASSI/LTR/TRIM
consensus size: 13912bp; fragments: 29738; full length: 13 (>=12520.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

