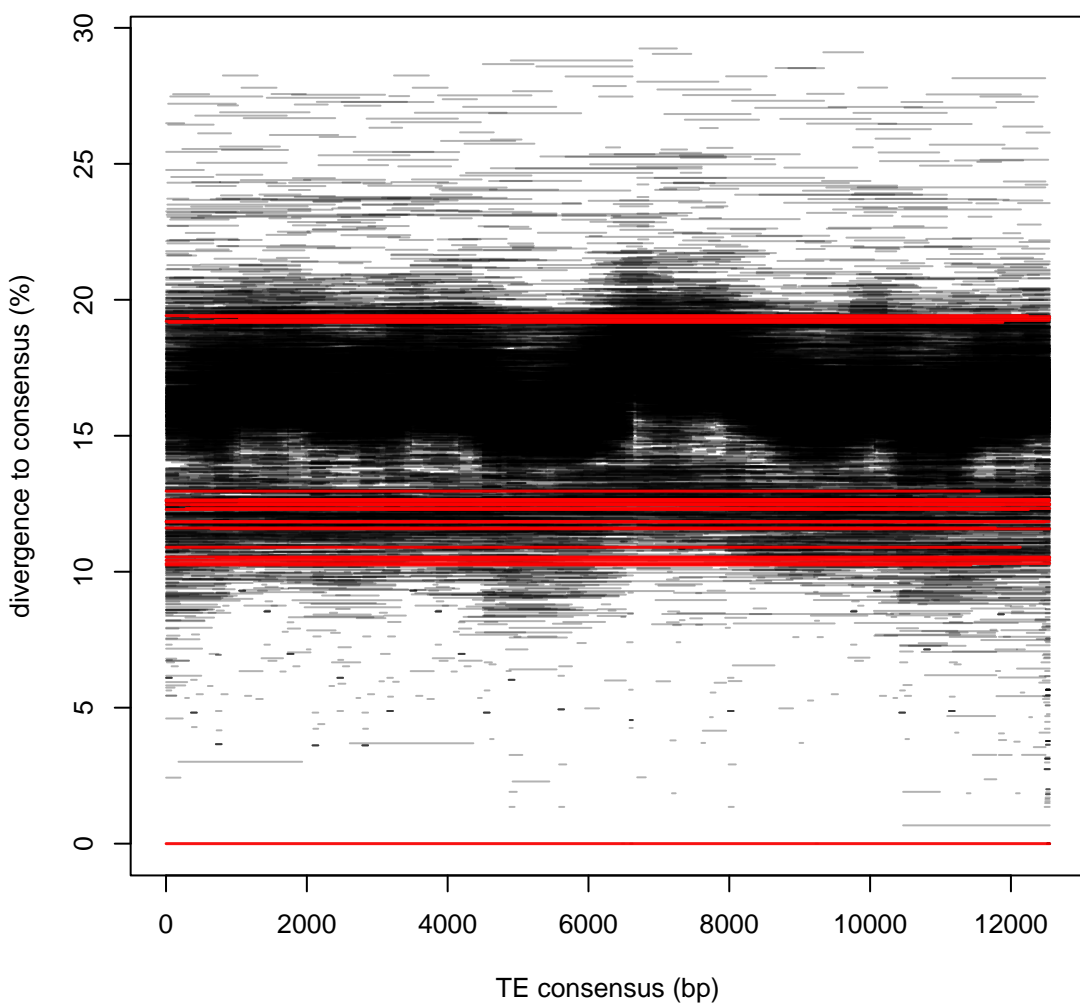
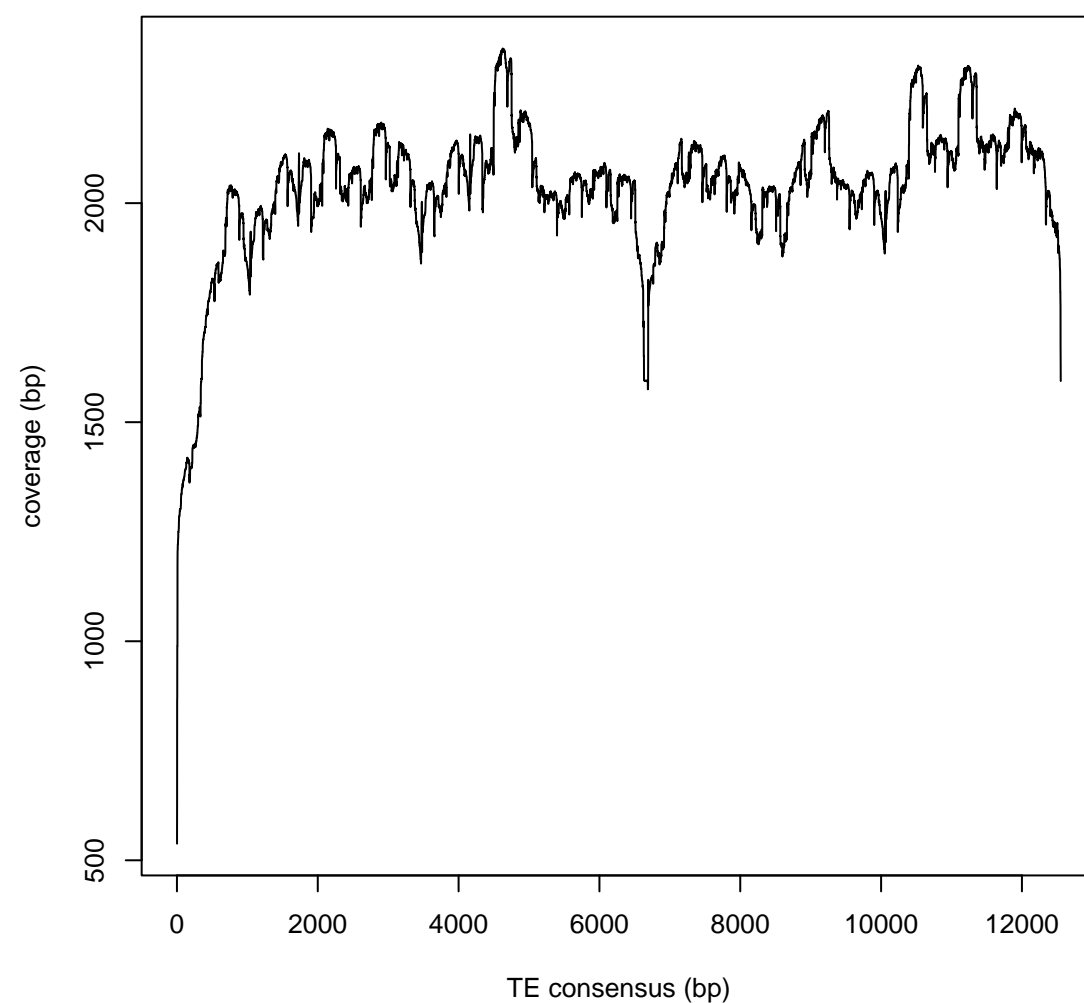


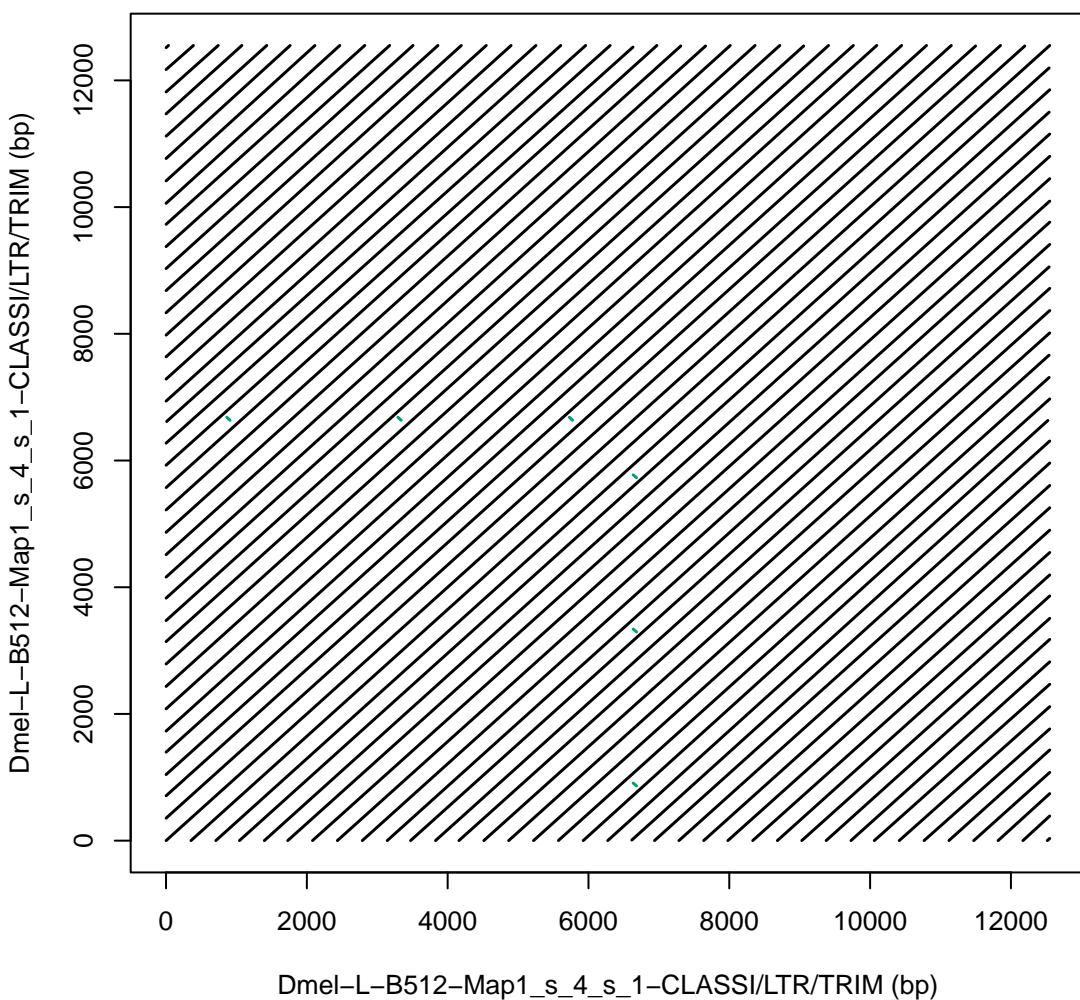
TE: Dmel-L-B512-Map1_s_4_s_1-CLASSI/LTR/TRIM
consensus size: 12551bp; fragments: 23290; full length: 27 (≥ 11295.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

