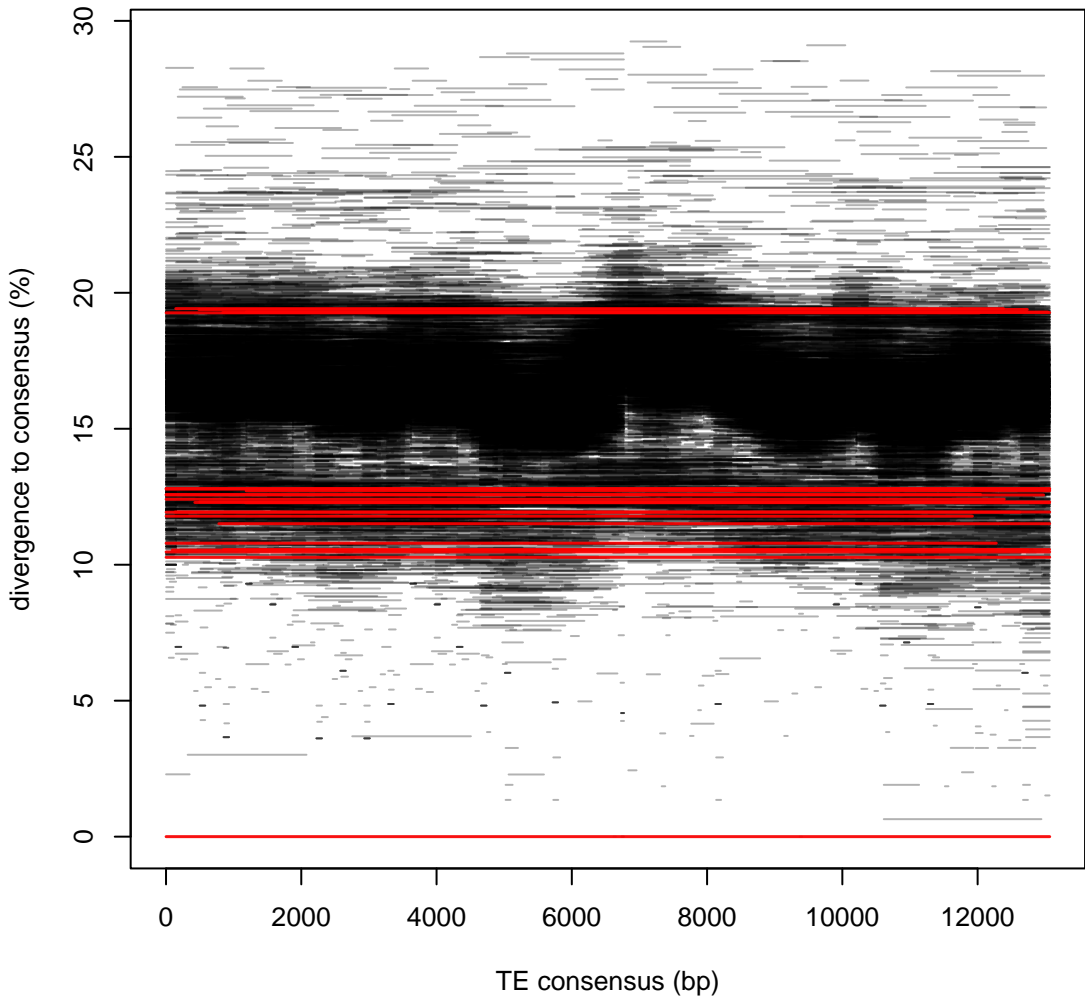
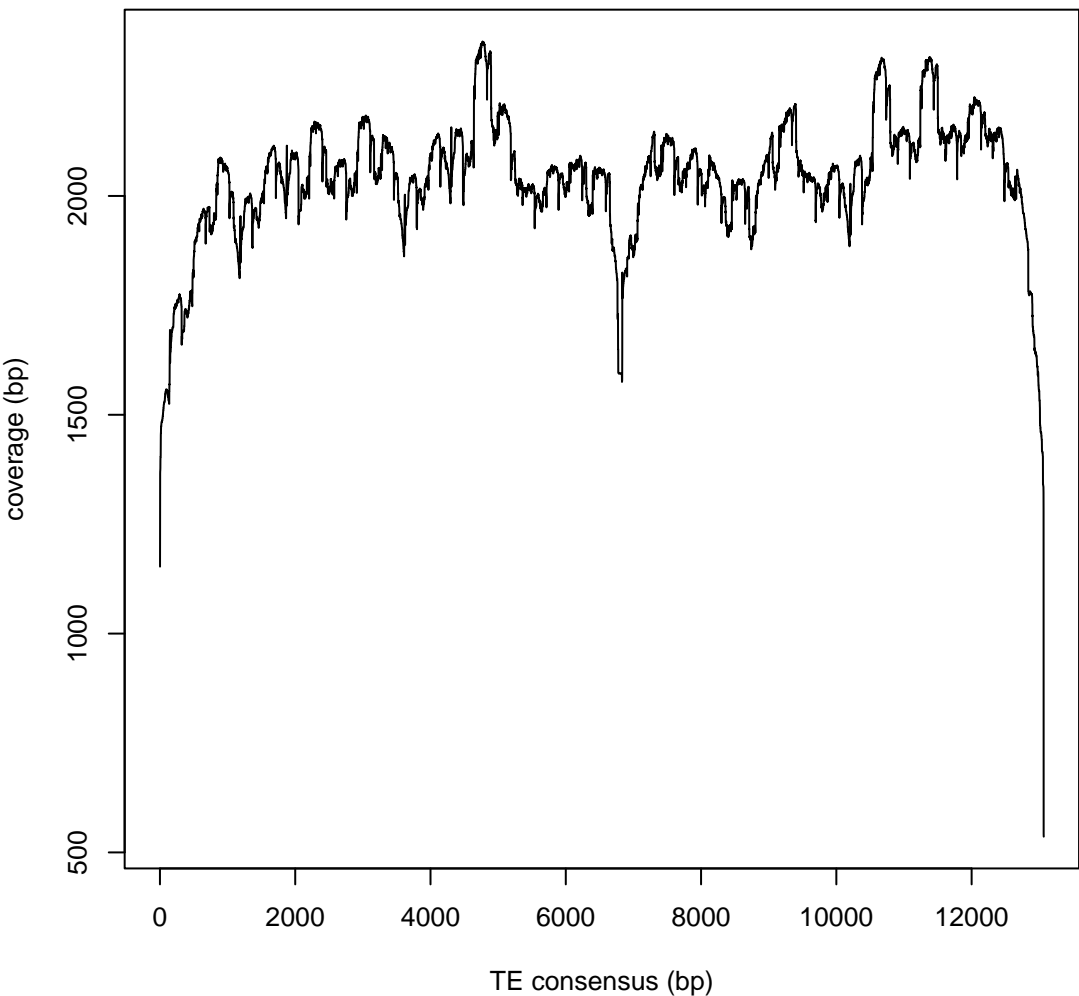


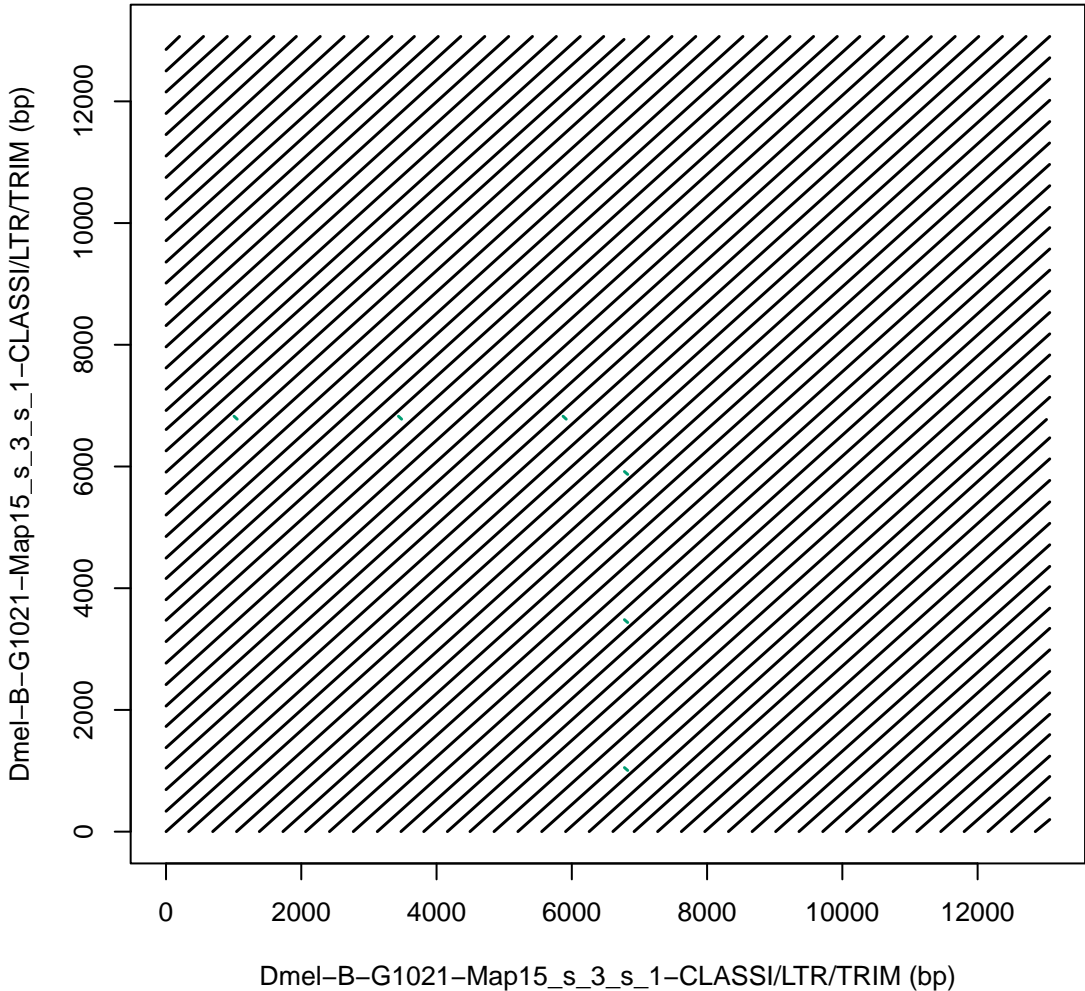
TE: Dmel-B-G1021-Map15_s_3_s_1-CLASSI/LTR/TRIM
consensus size: 13066bp; fragments: 24076; full length: 24 (>=11759.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

