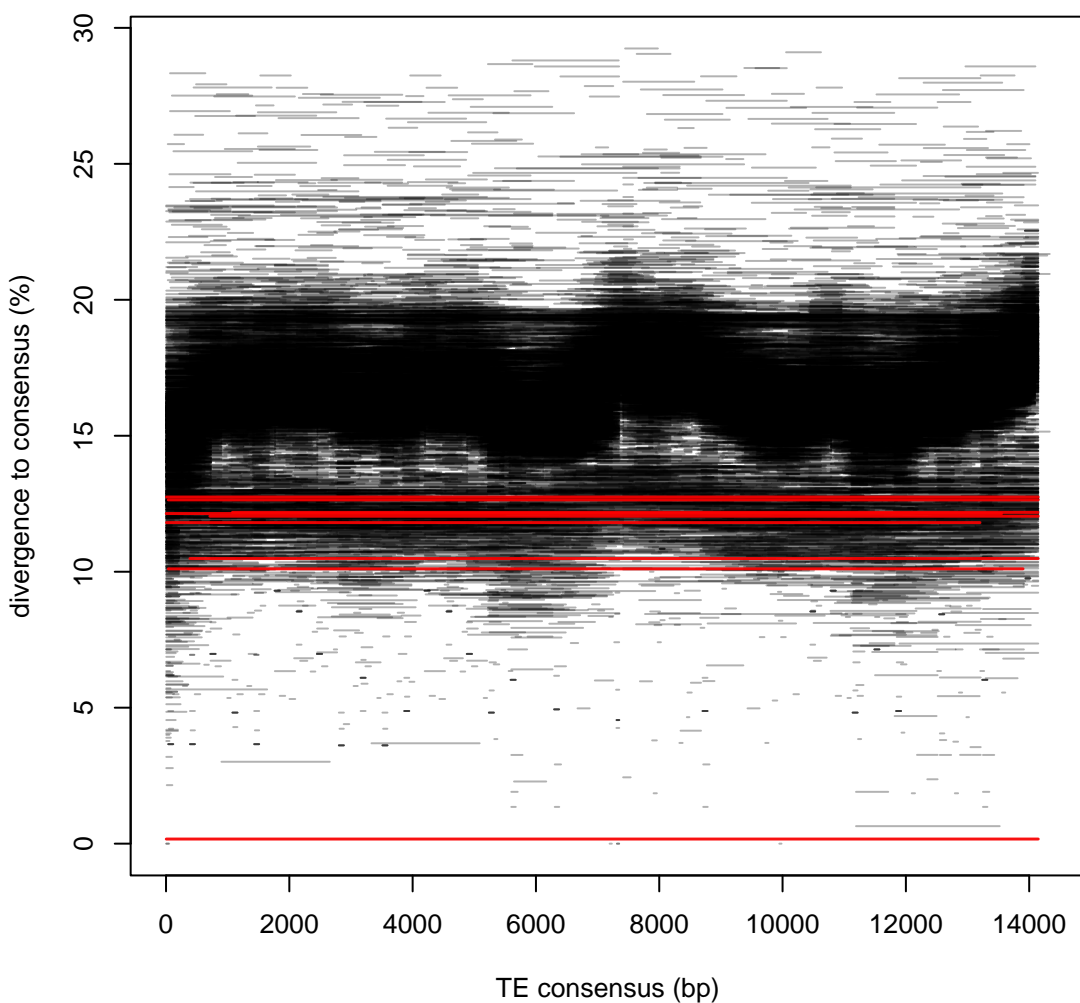
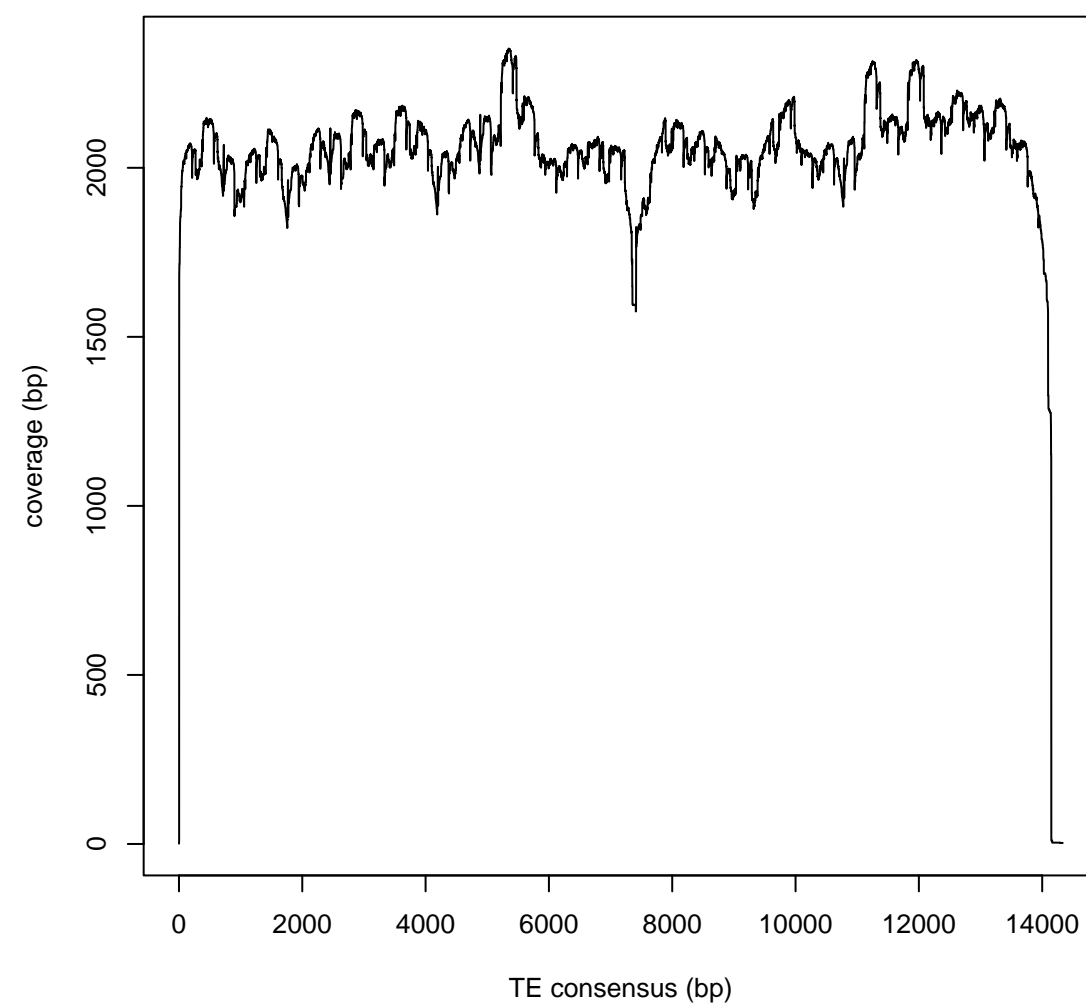


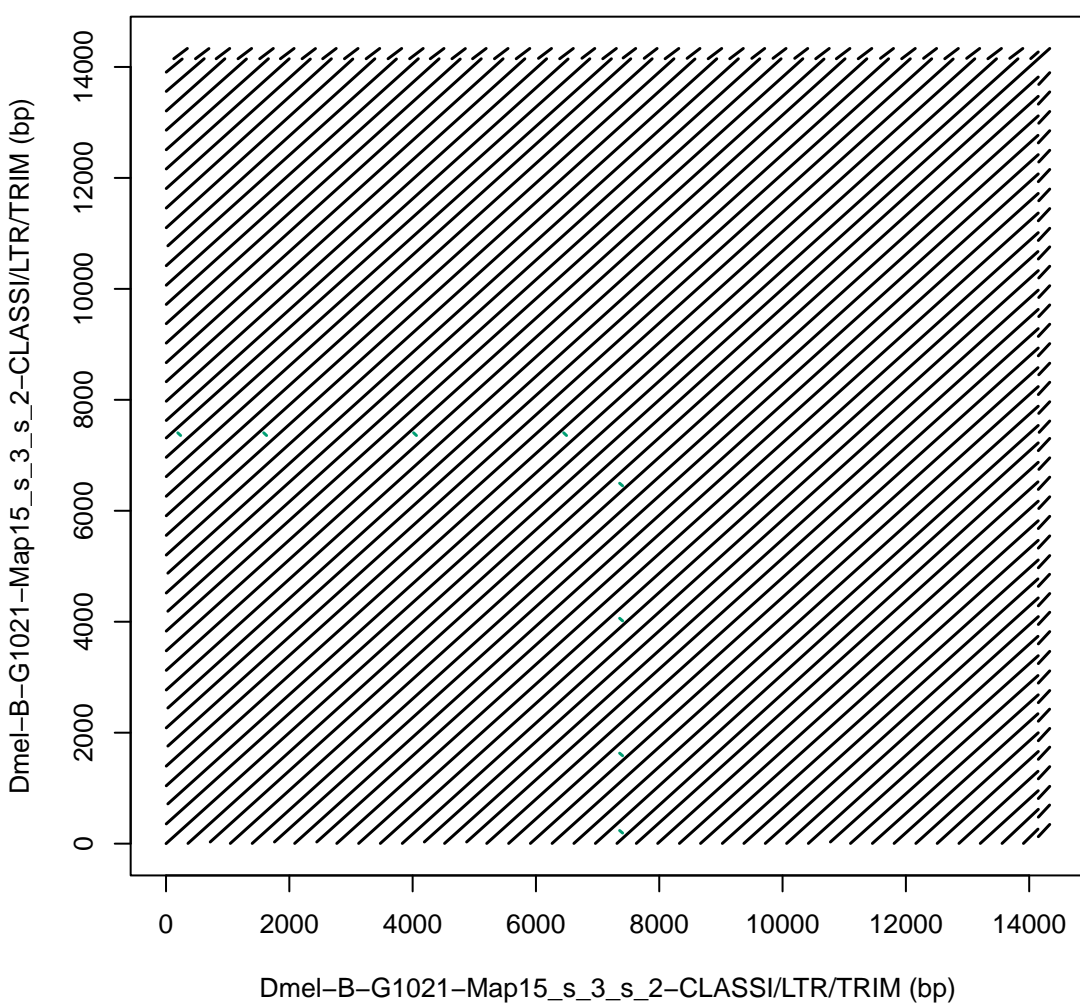
TE: Dmel-B-G1021-Map15\_s\_3\_s\_2-CLASSI/LTR/TRIM  
consensus size: 14333bp; fragments: 26477; full length: 9 (>=12899.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

