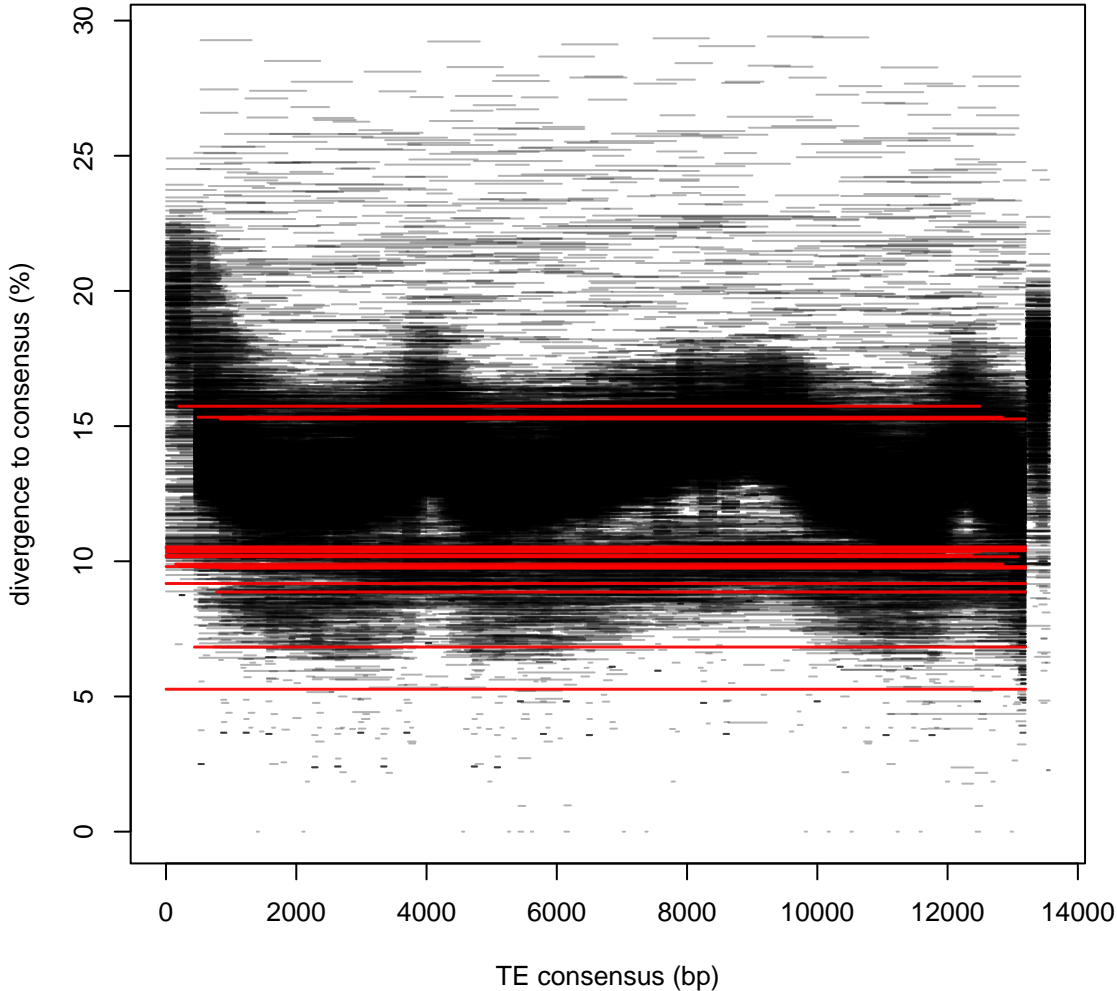
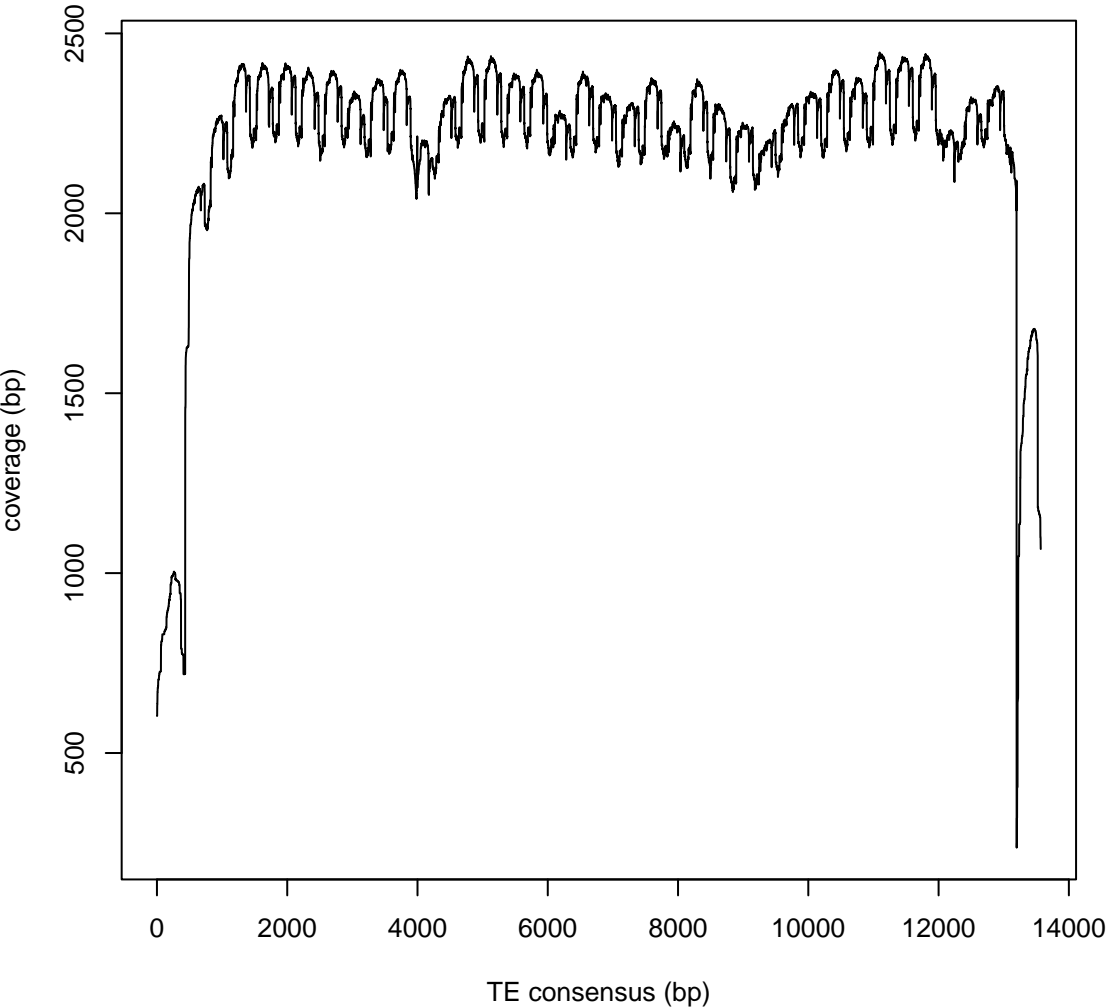


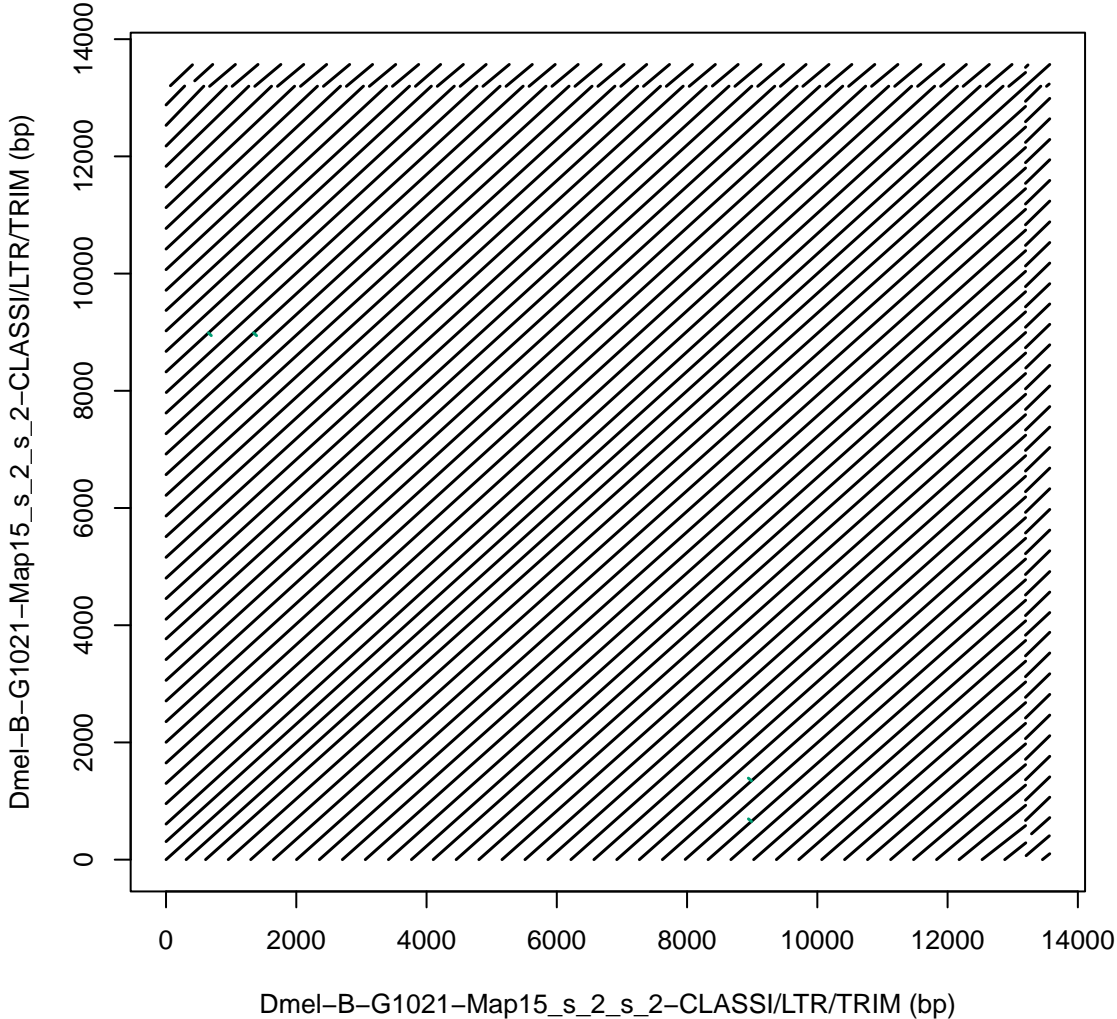
TE: Dmel-B-G1021-Map15\_s\_2\_s\_2-CLASSI/LTR/TRIM  
consensus size: 13568bp; fragments: 31916; full length: 18 (>=12211.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

