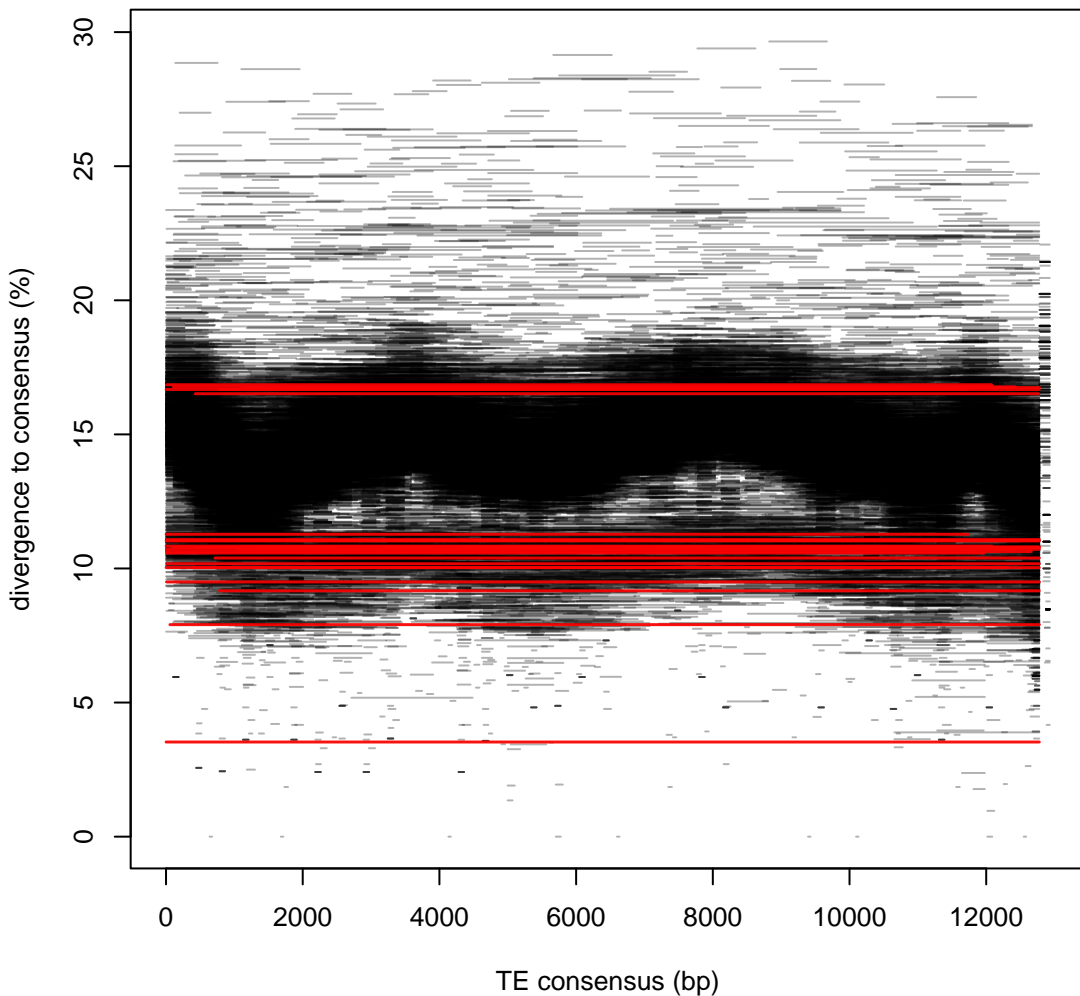
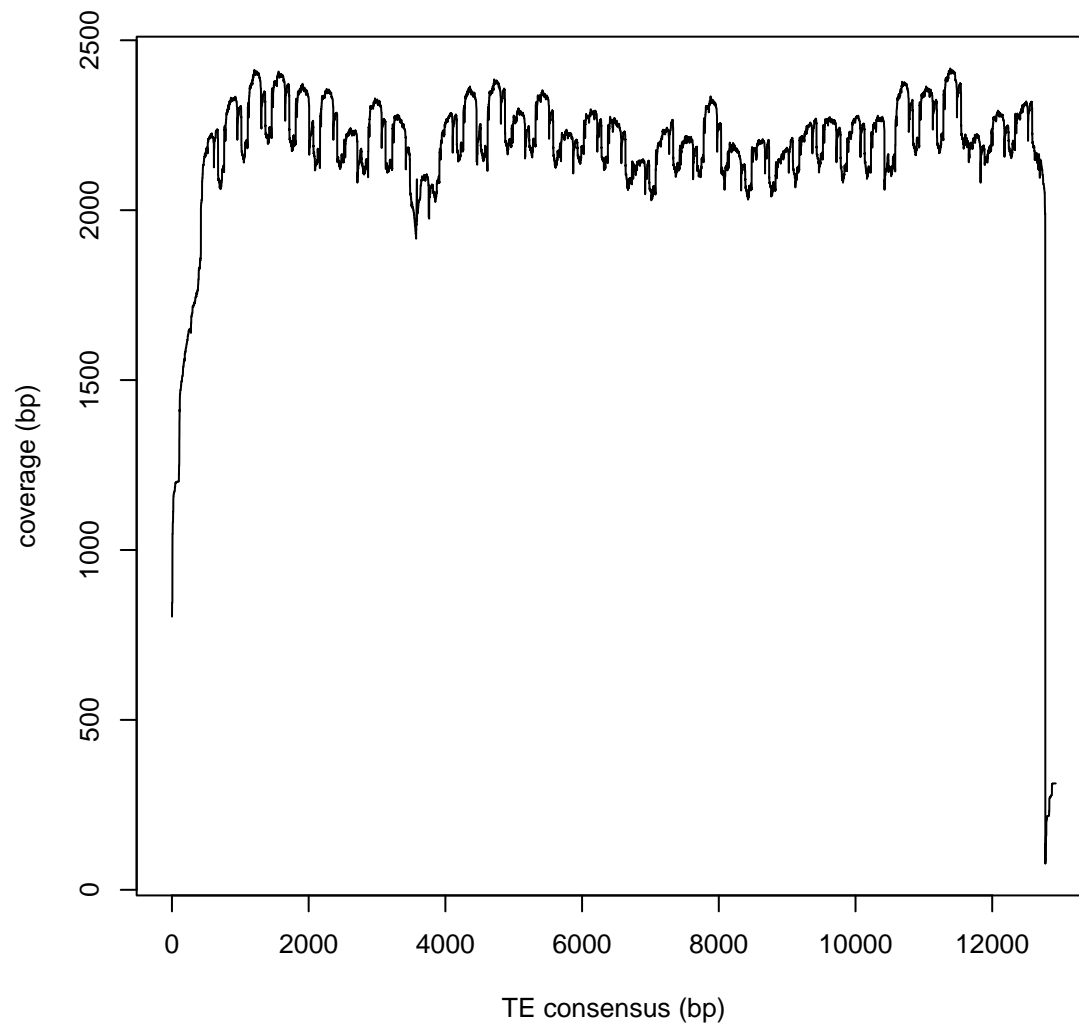


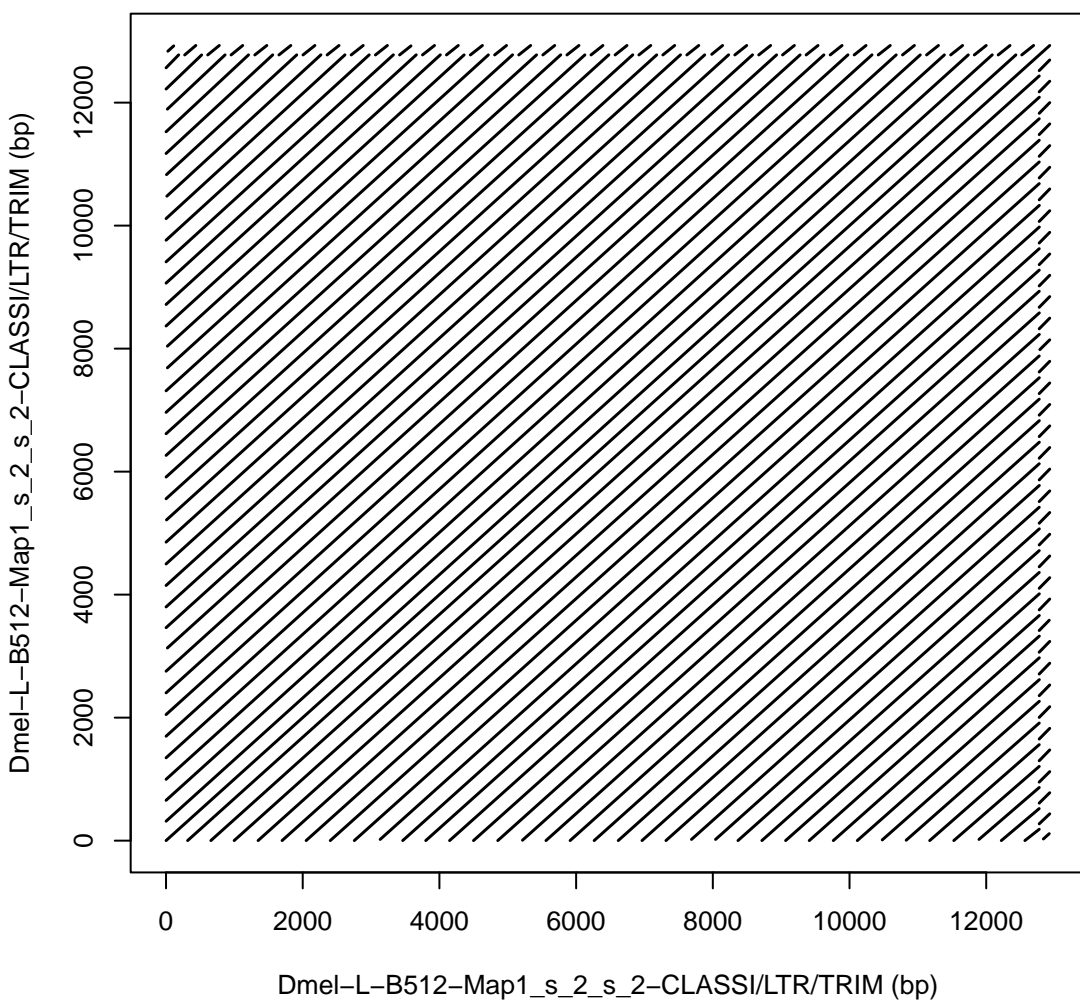
TE: Dmel-L-B512-Map1\_s\_2\_s\_2-CLASSI/LTR/TRIM  
consensus size: 12929bp; fragments: 28096; full length: 26 (>=11636.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

