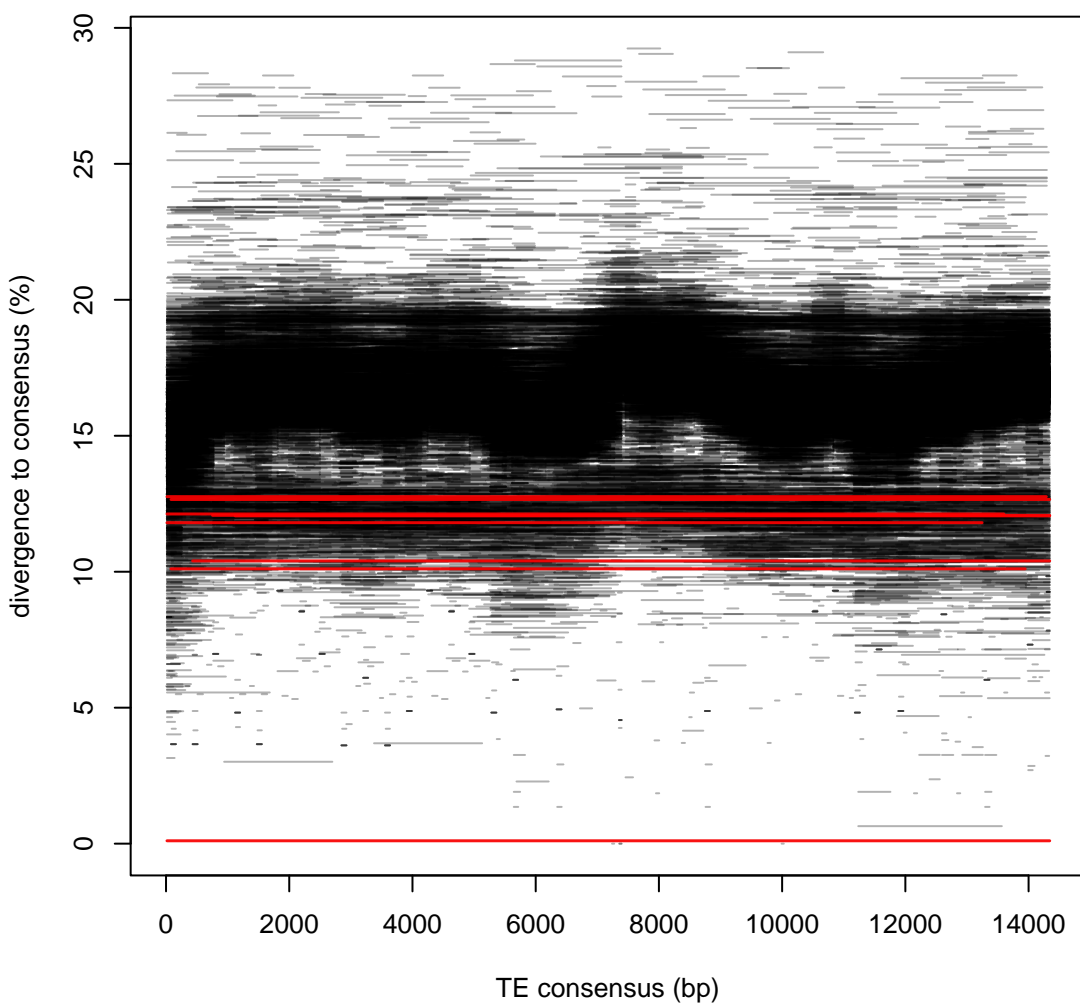
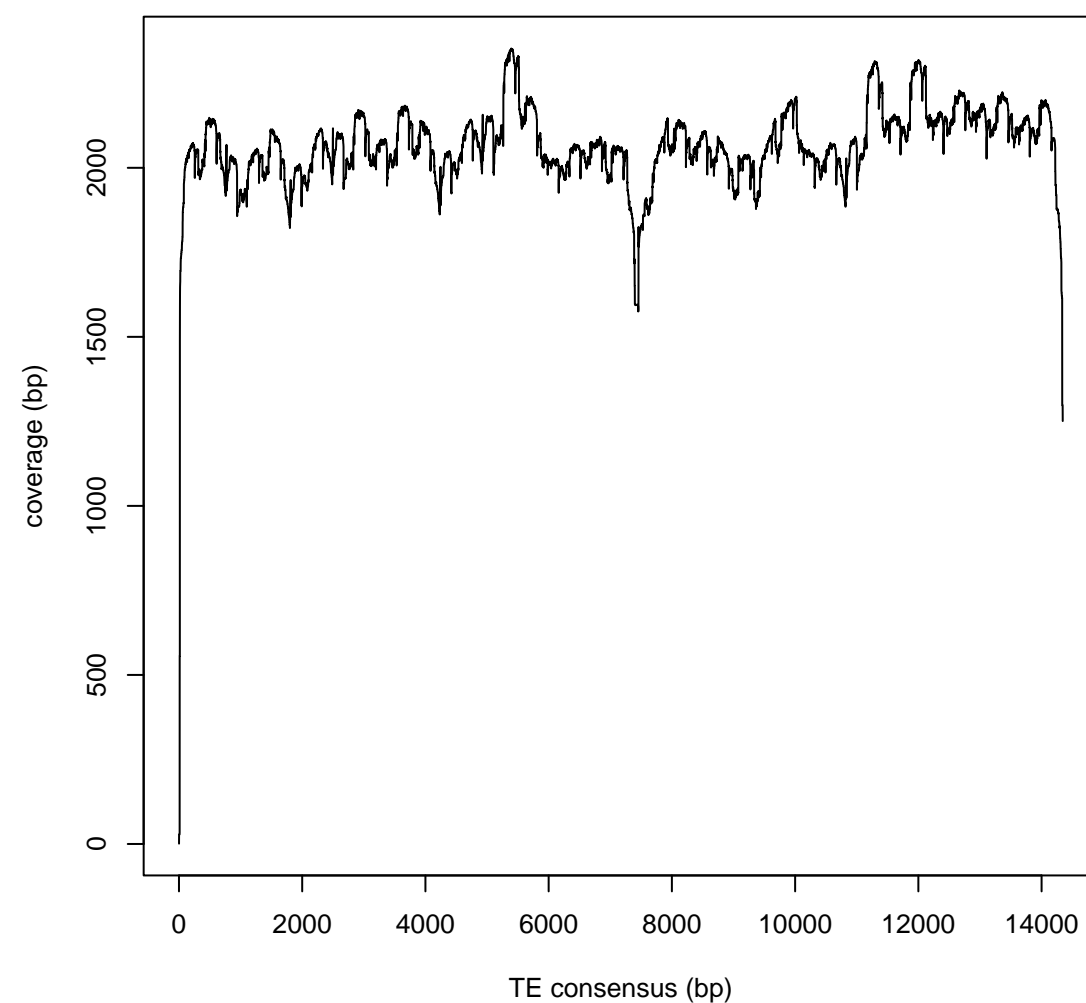


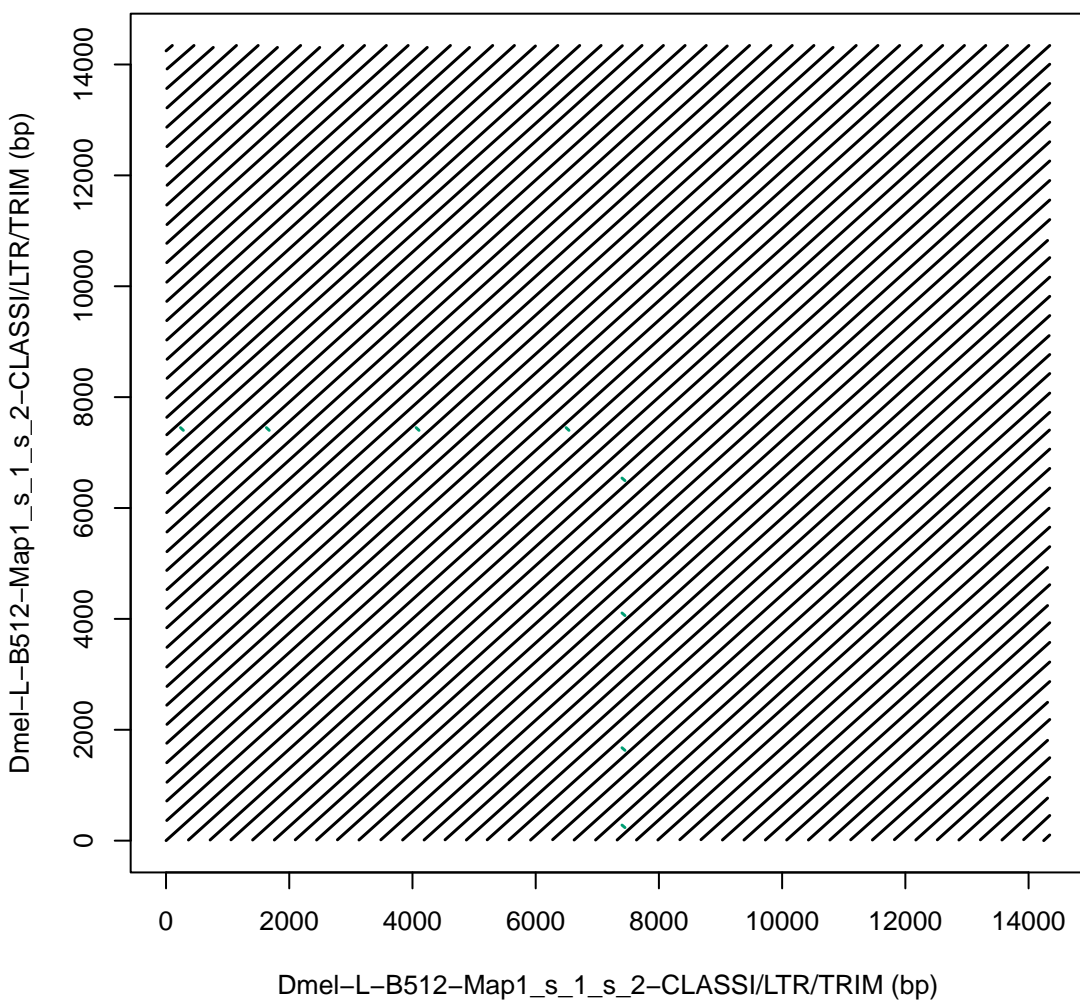
TE: Dmel-L-B512-Map1\_s\_1\_s\_2-CLASSI/LTR/TRIM  
consensus size: 14343bp; fragments: 27049; full length: 9 (>=12908.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

