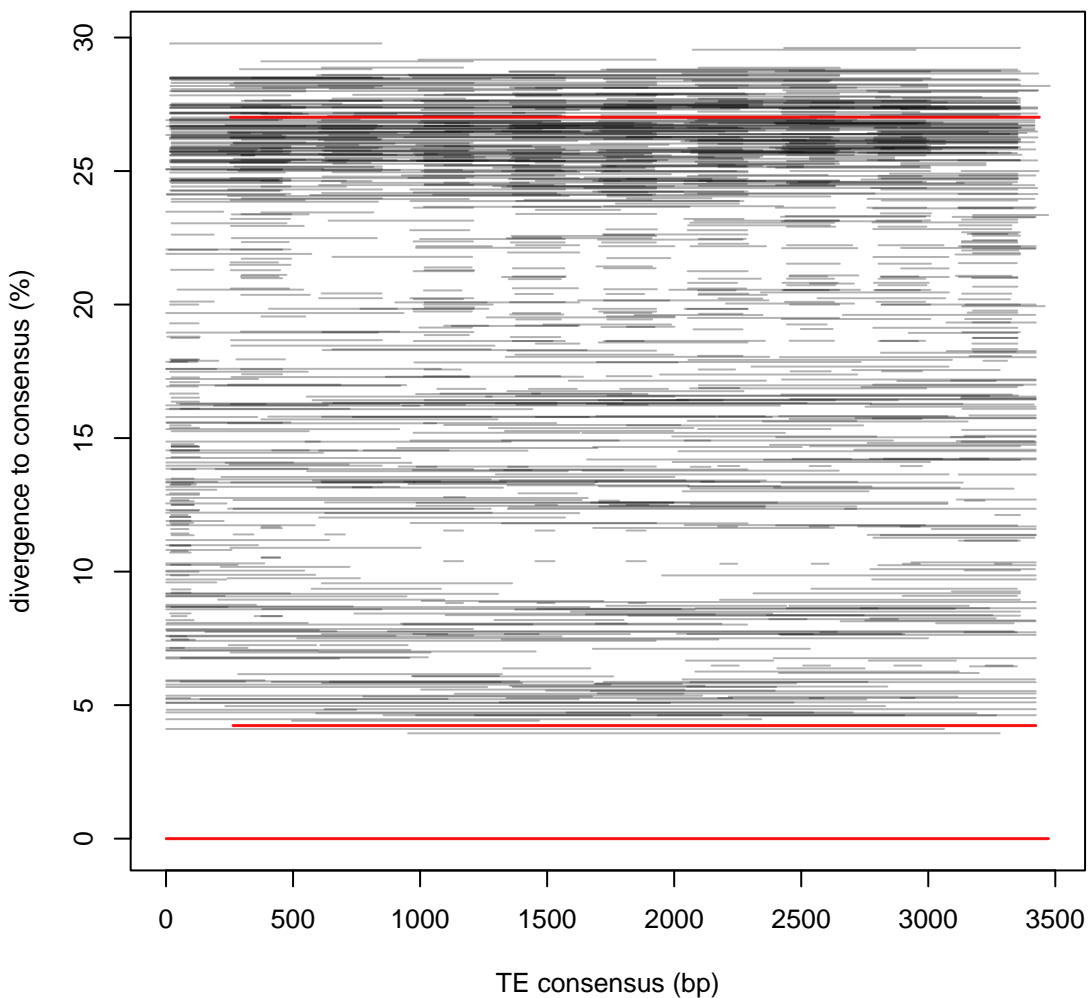
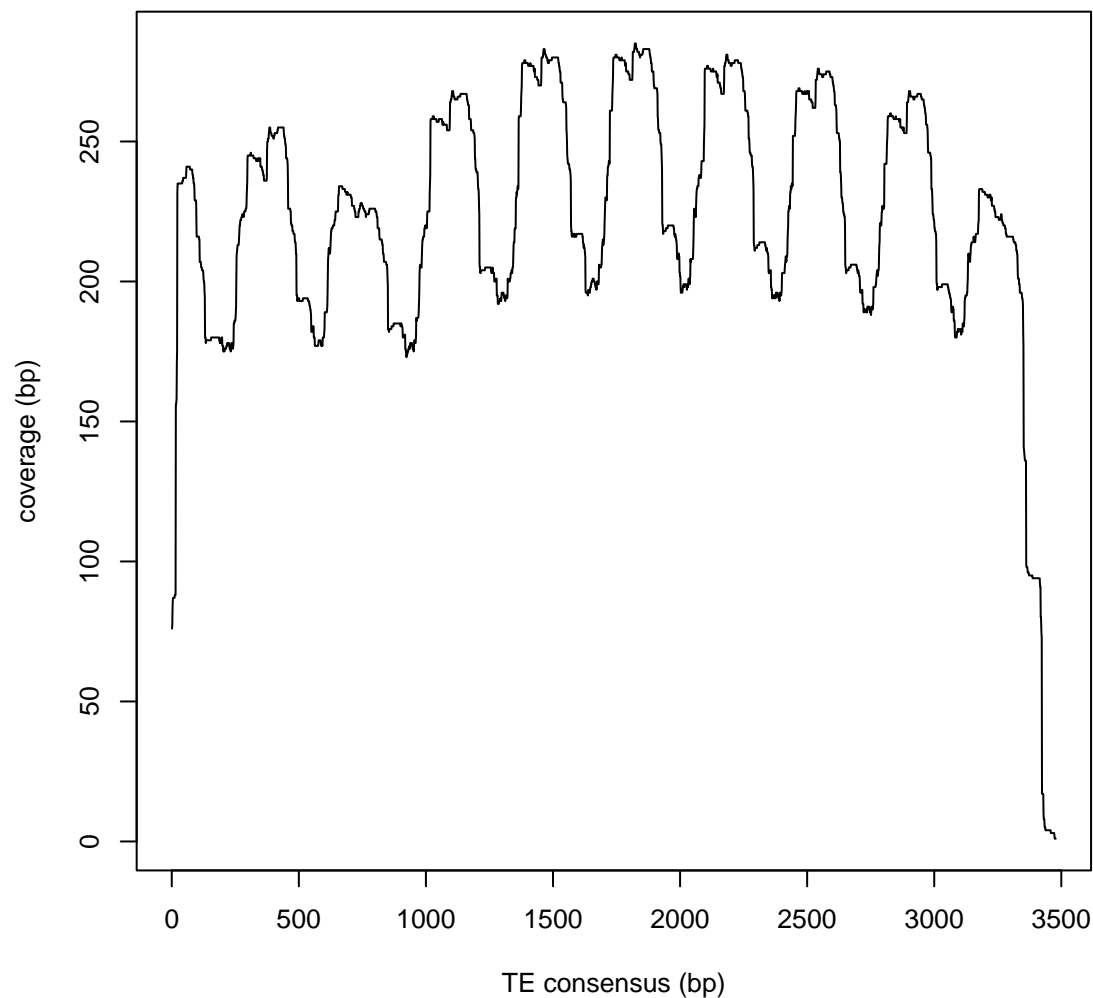


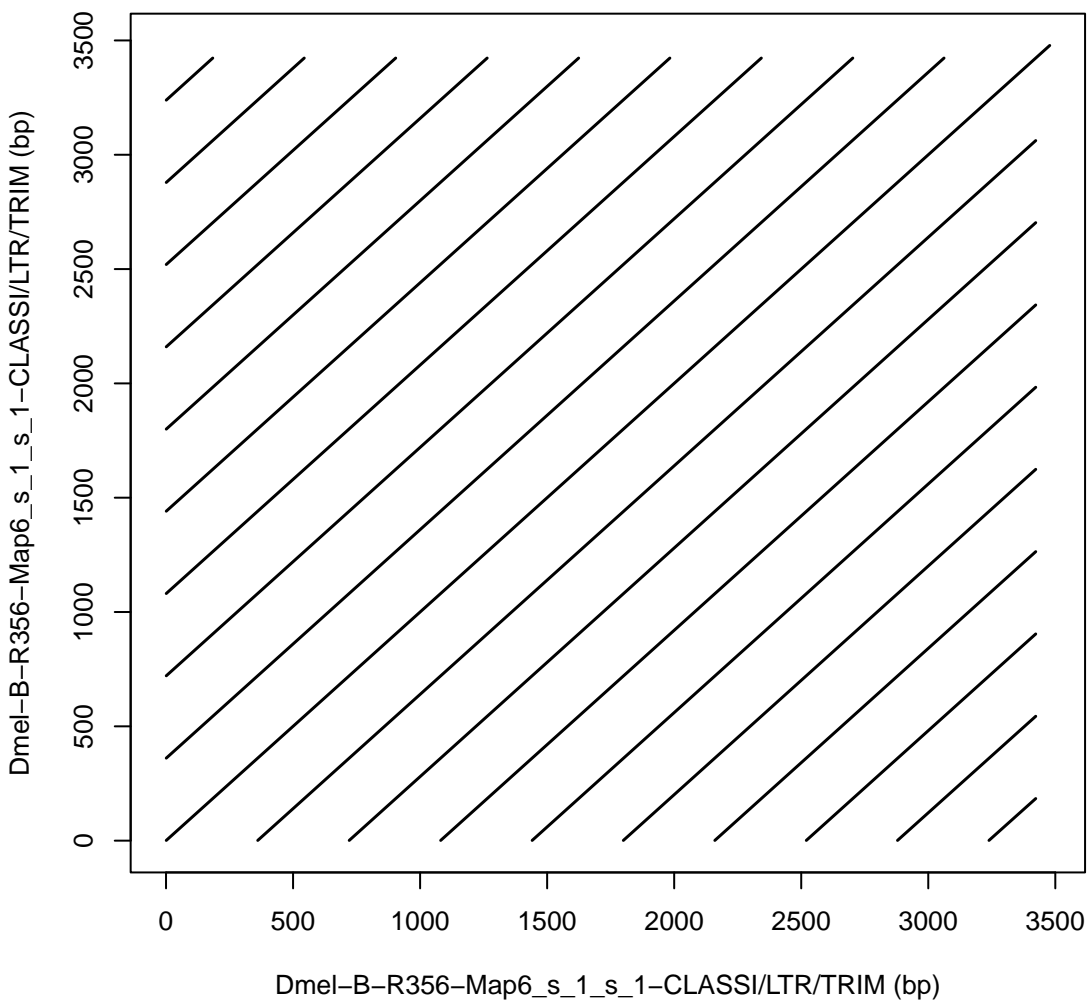
TE: Dmel-B-R356-Map6\_s\_1\_s\_1-CLASSI/LTR/TRIM  
consensus size: 3478bp; fragments: 1222; full length: 3 (>=3130.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

