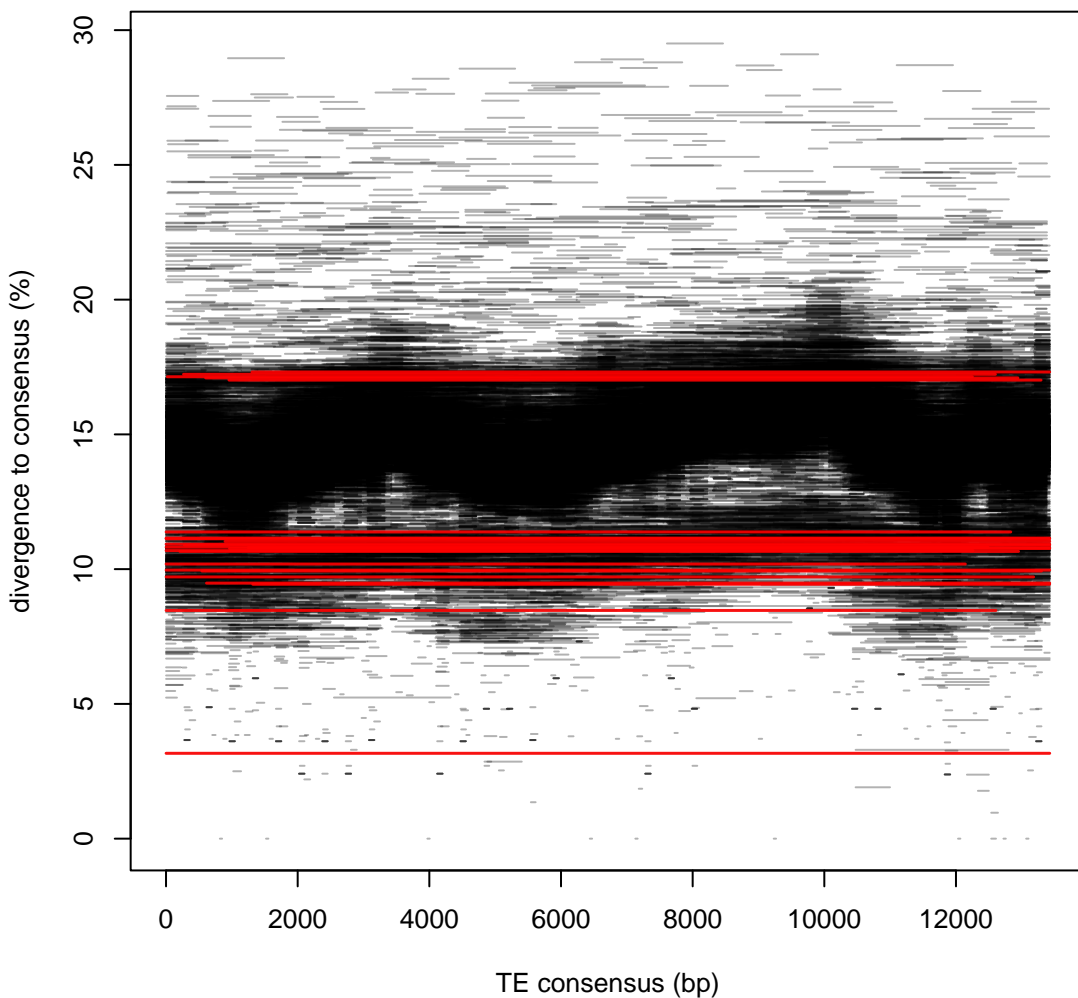
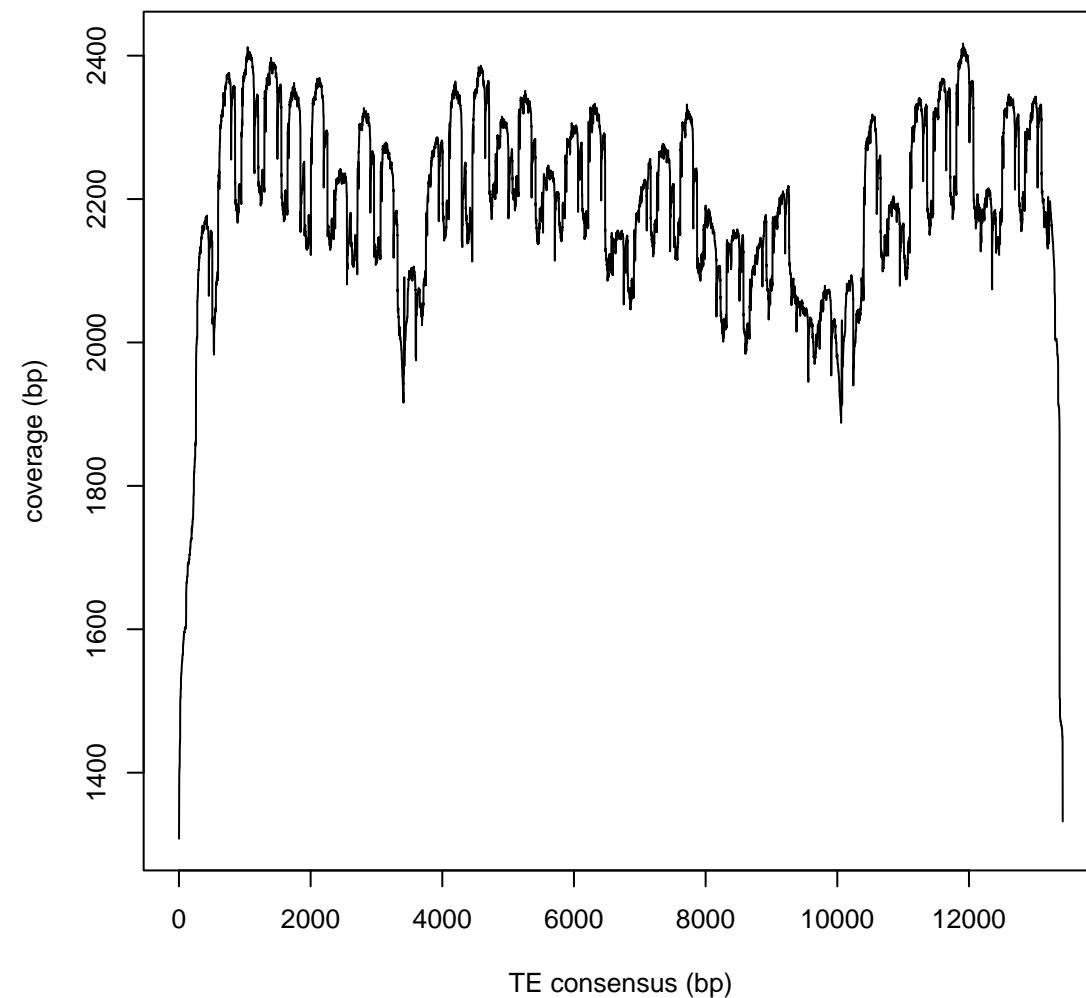


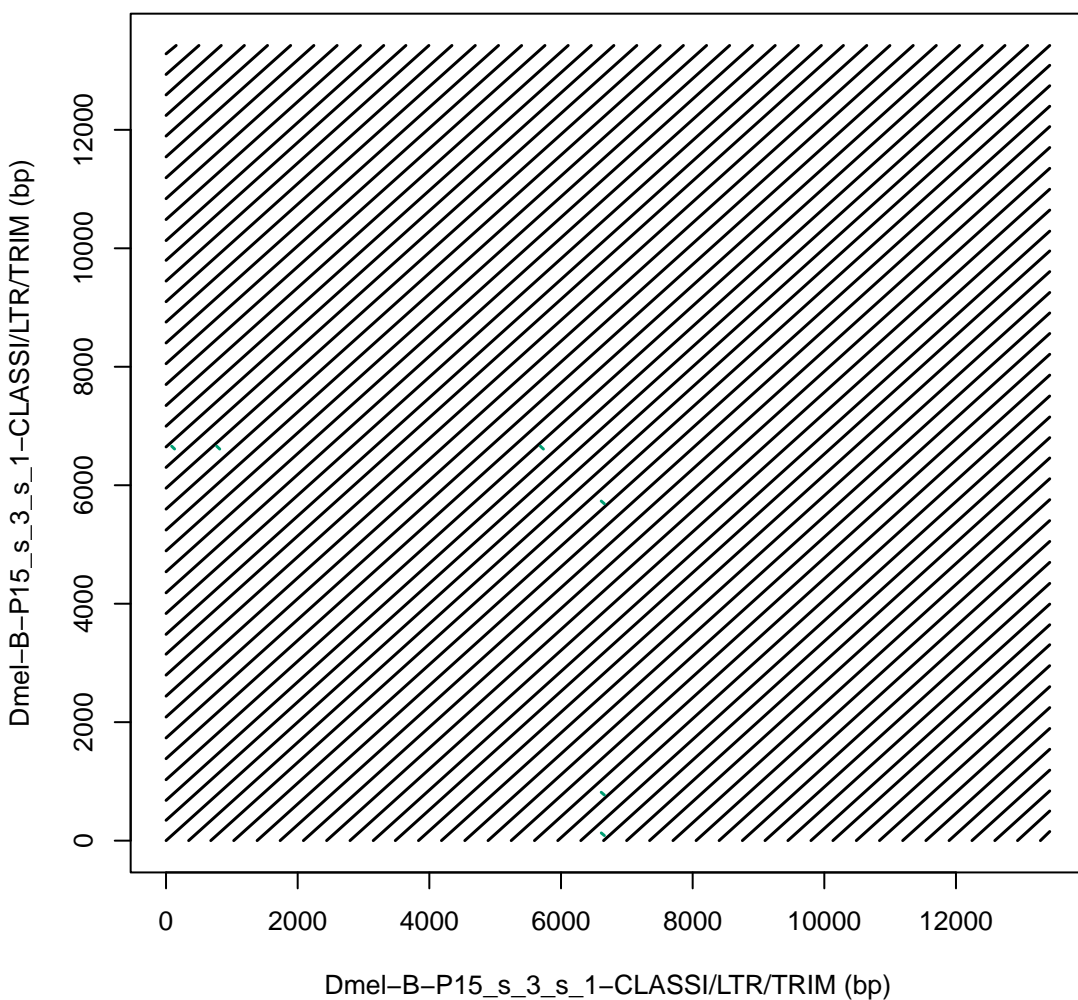
TE: Dmel-B-P15\_s\_3\_s\_1-CLASSI/LTR/TRIM  
consensus size: 13424bp; fragments: 28758; full length: 24 ( $\geq 12081.6$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

