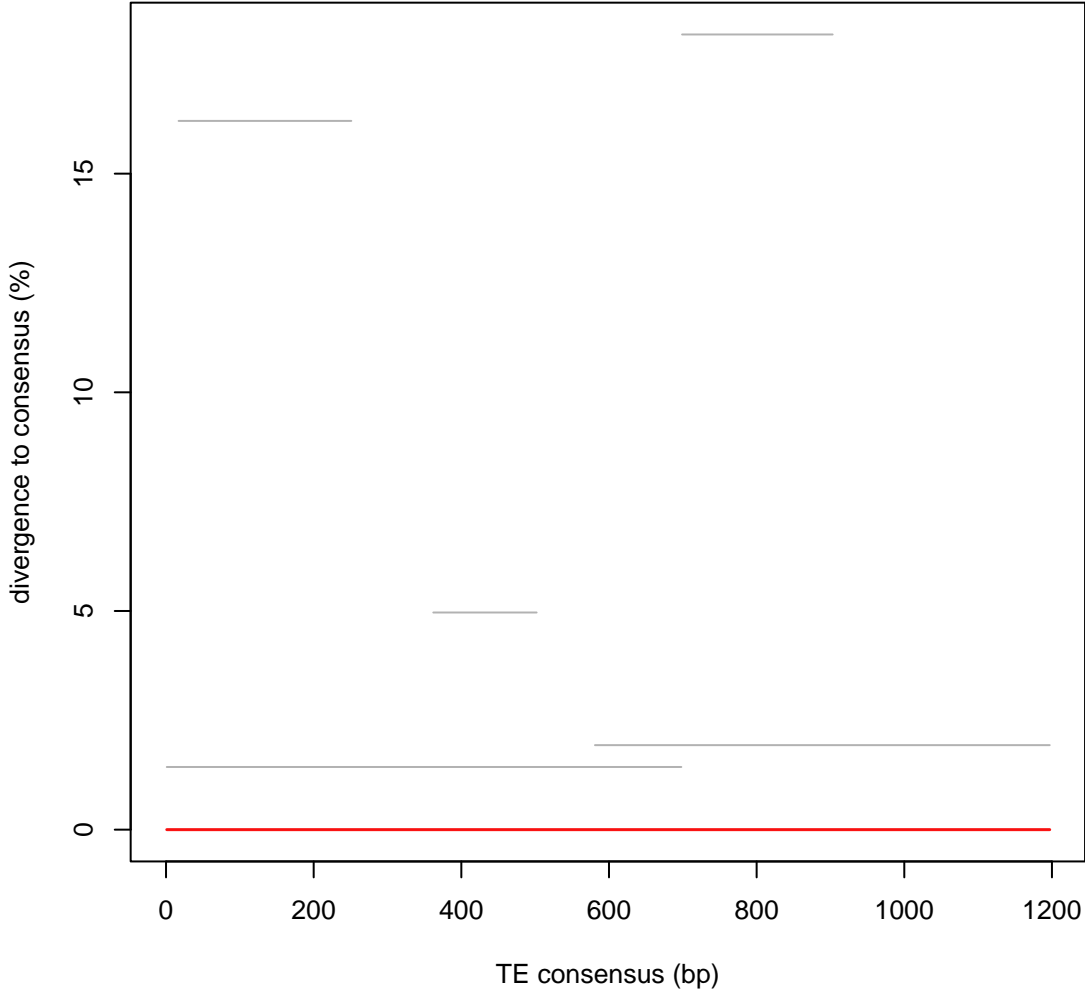
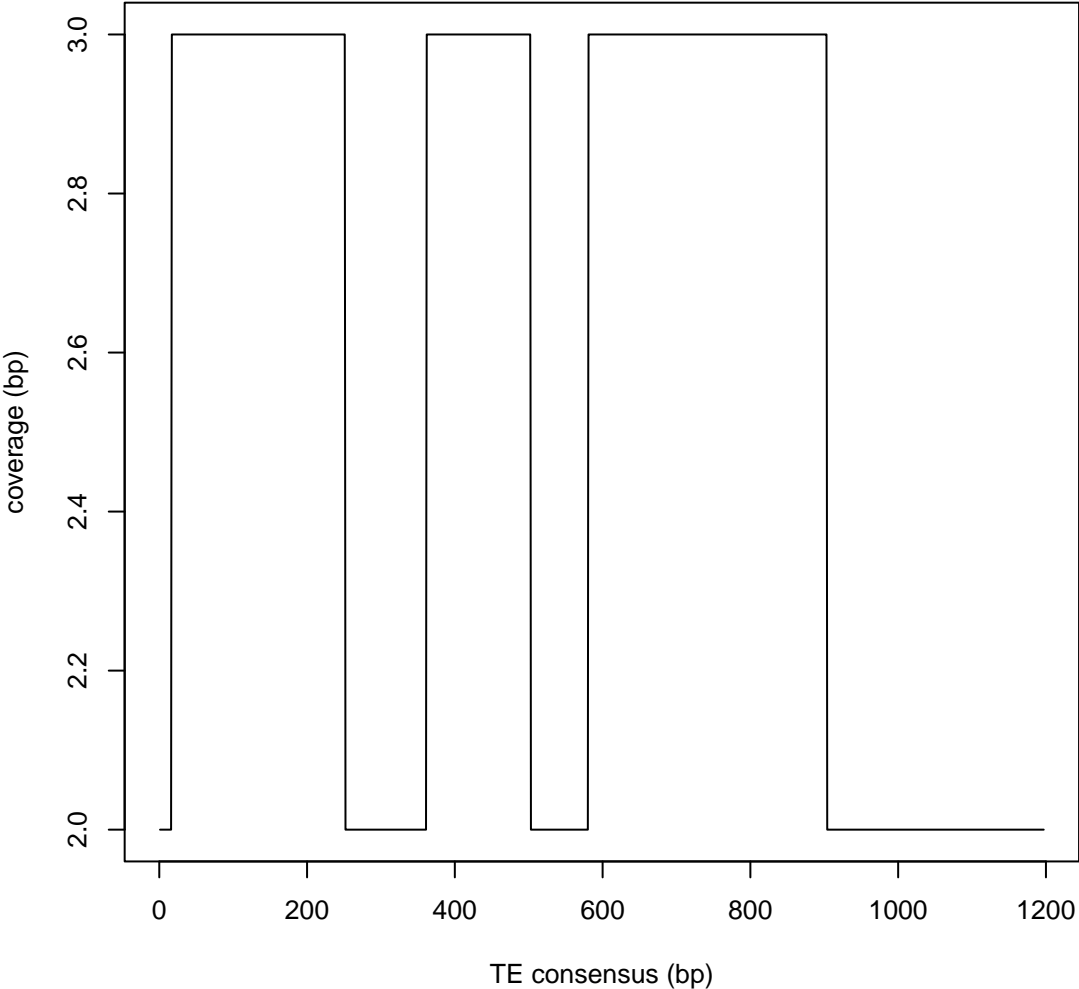


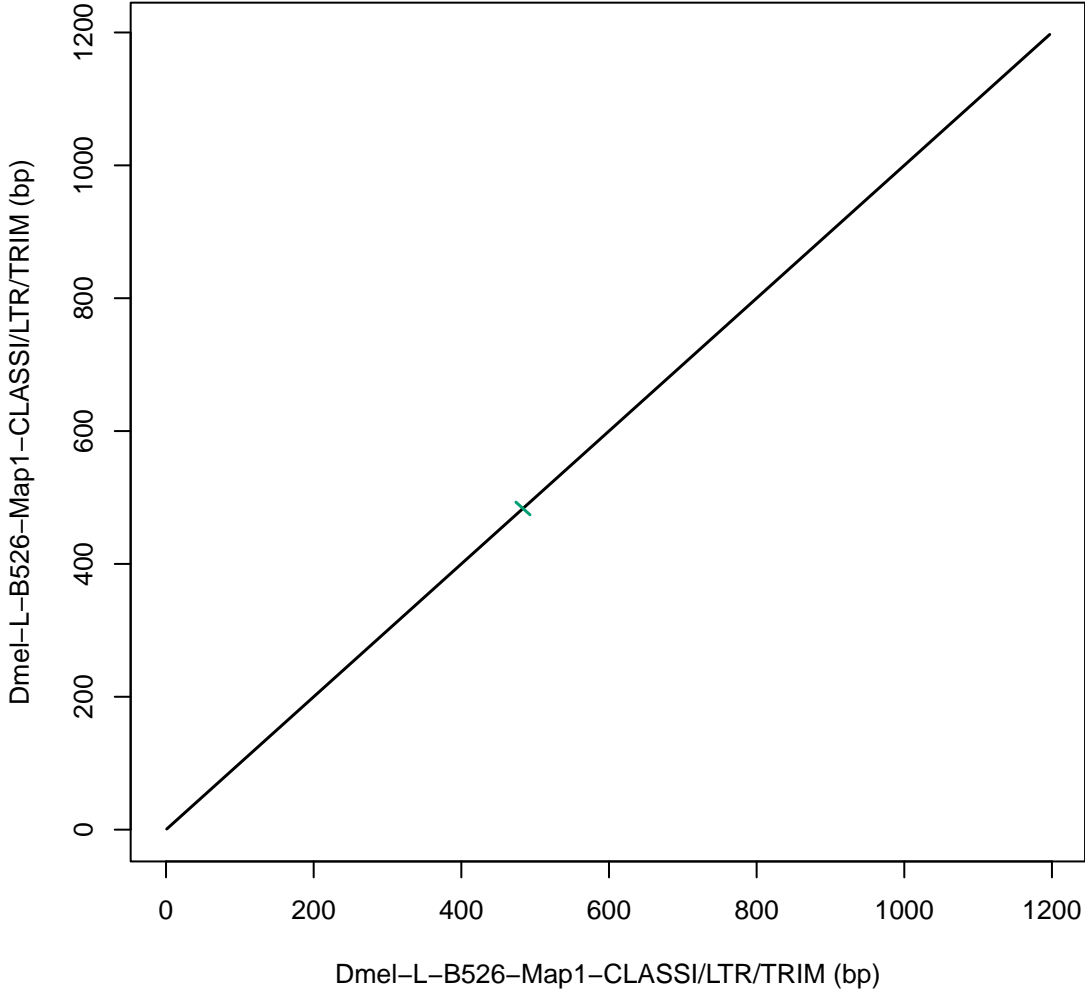
TE: Dmel-L-B526-Map1-CLASSI/LTR/TRIM  
consensus size: 1197bp; fragments: 6; full length: 1 (>=1077.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

