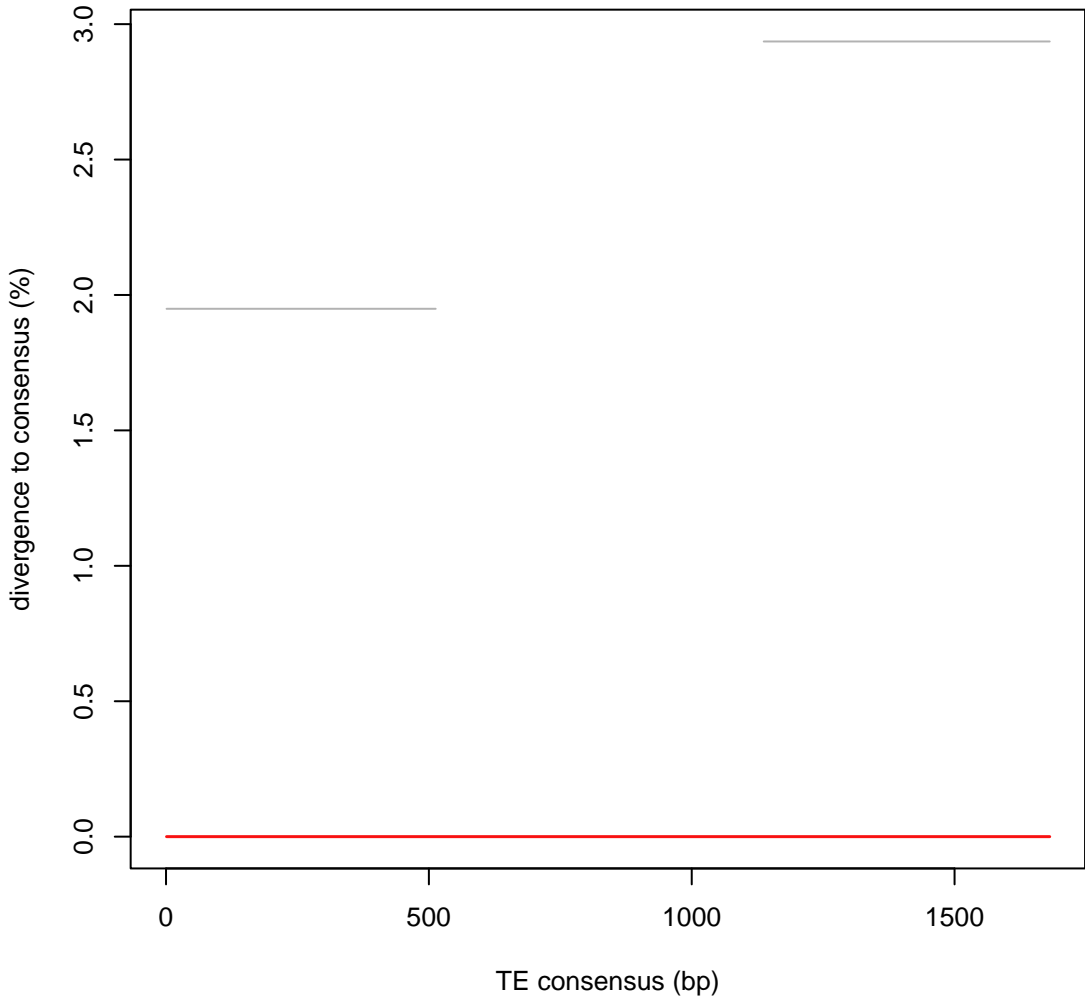
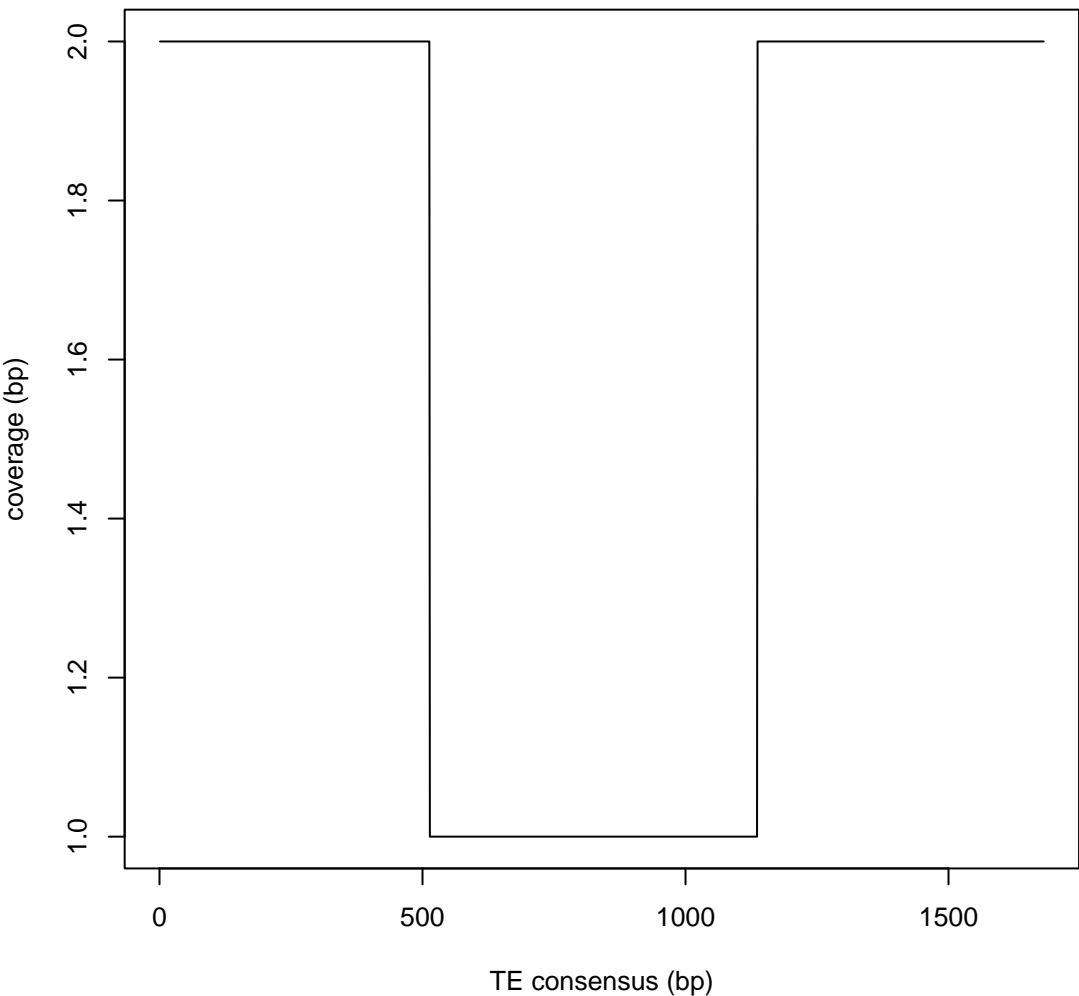


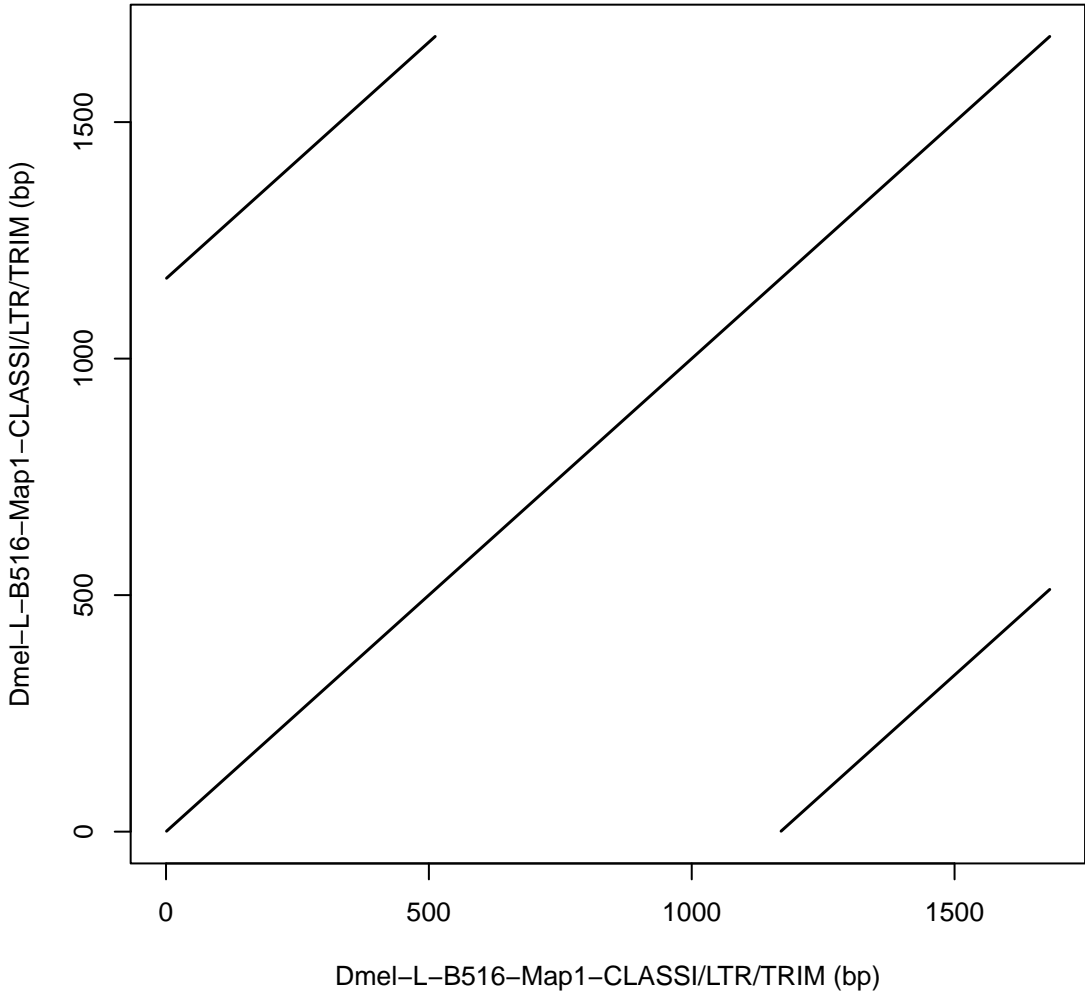
TE: Dmel-L-B516-Map1-CLASSI/LTR/TRIM  
consensus size: 1681bp; fragments: 3; full length: 1 (>=1512.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

