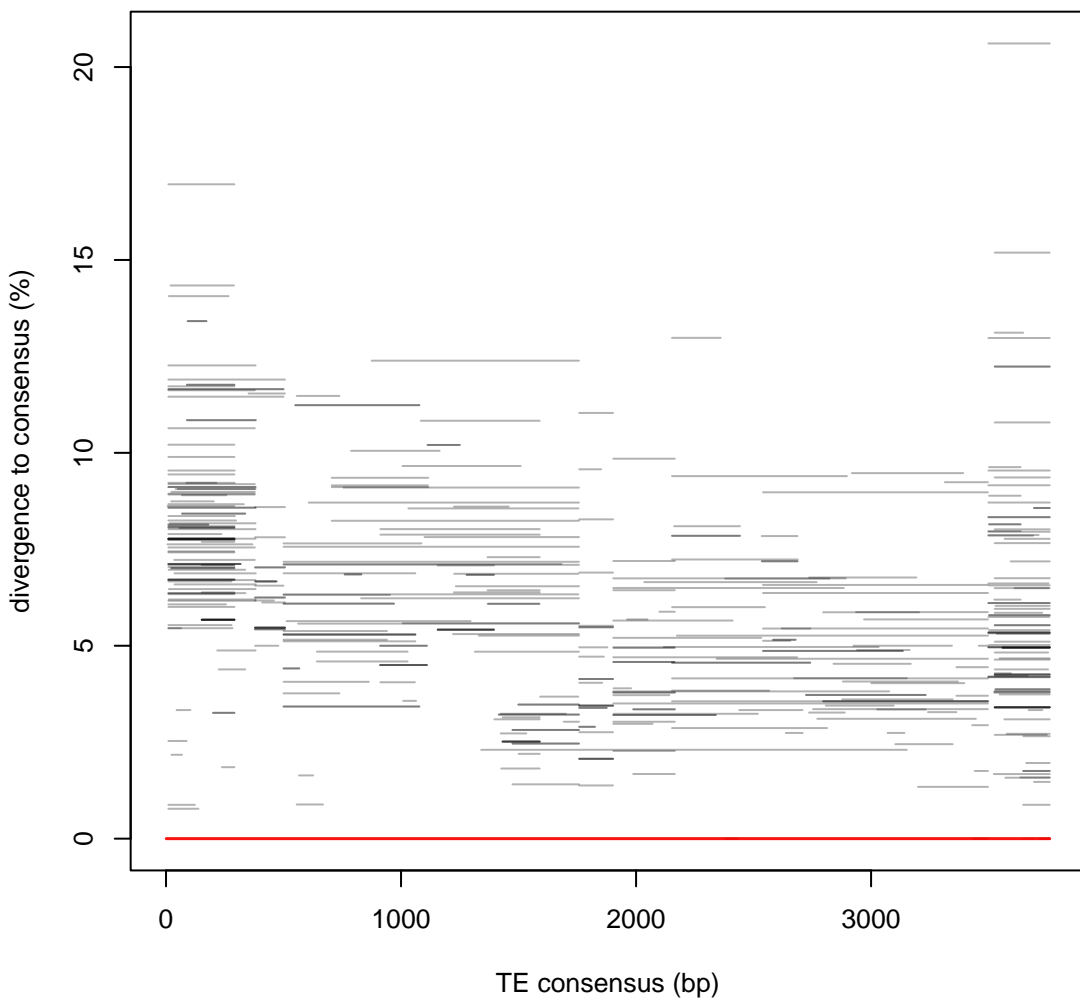
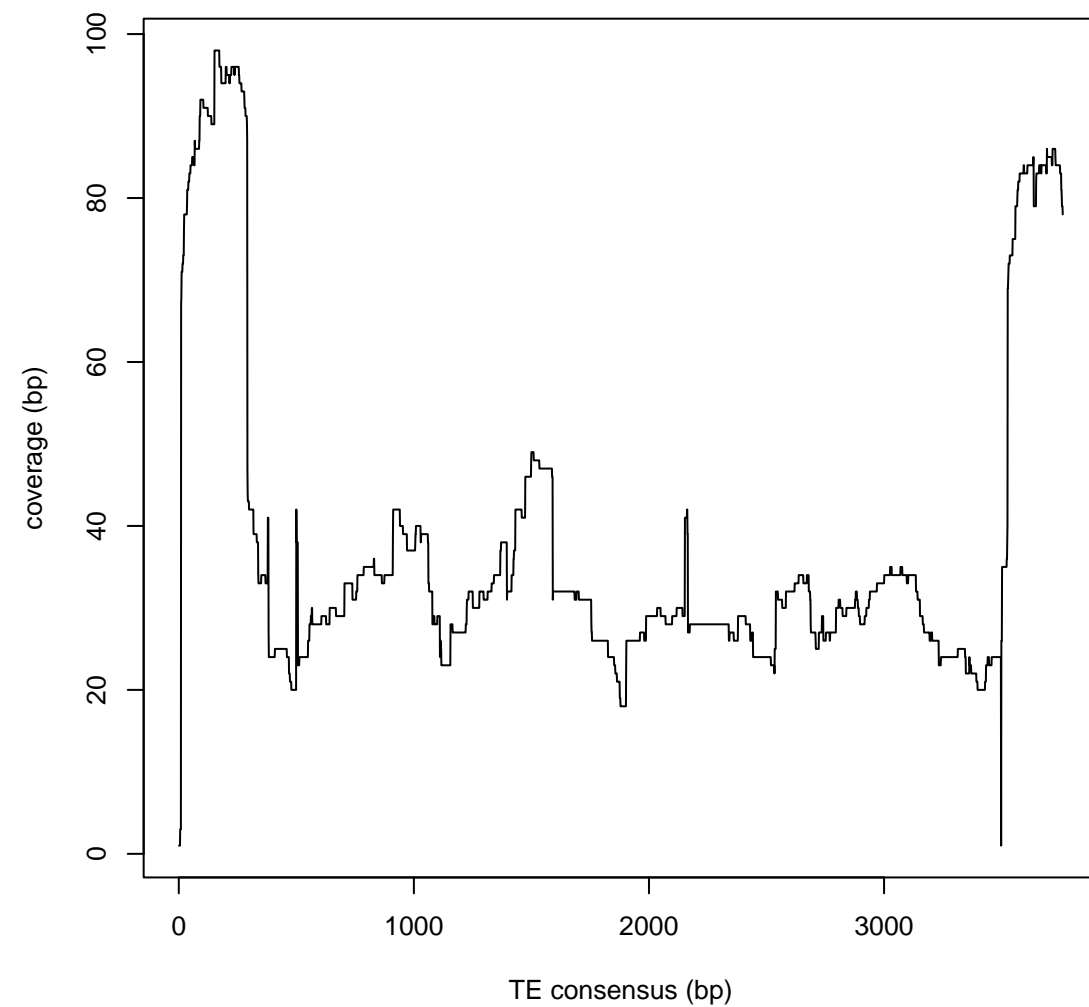


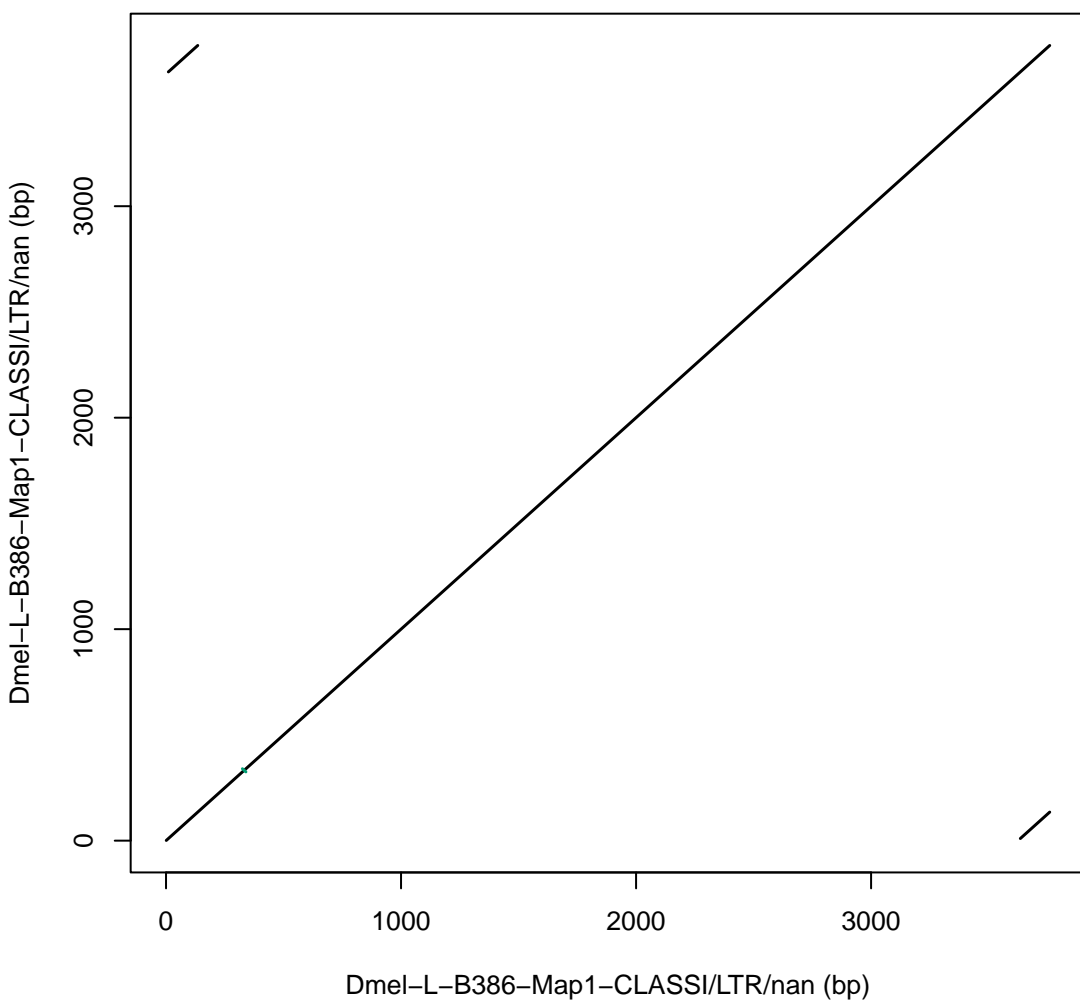
TE: Dmel-L-B386-Map1-CLASSI/LTR/nan
consensus size: 3760bp; fragments: 445; full length: 1 (>=3384bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

