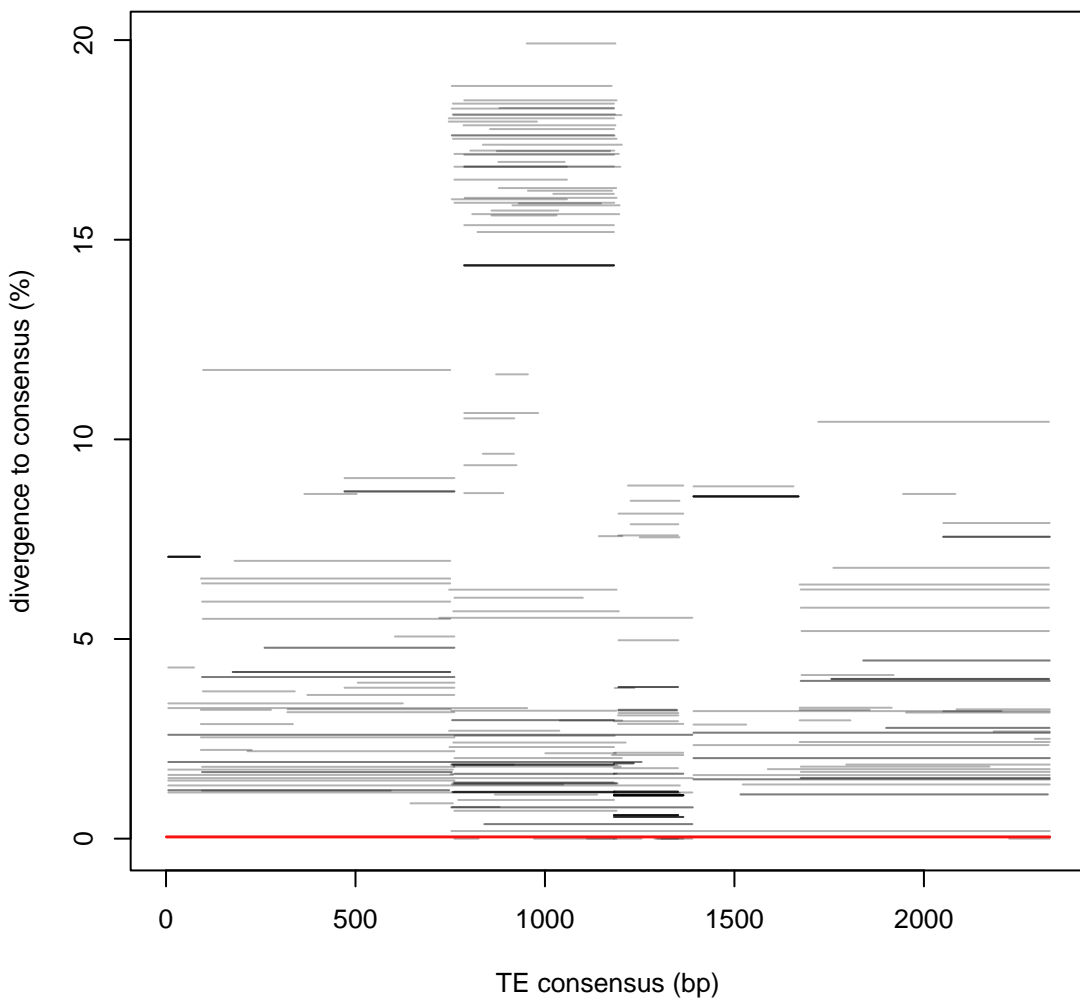
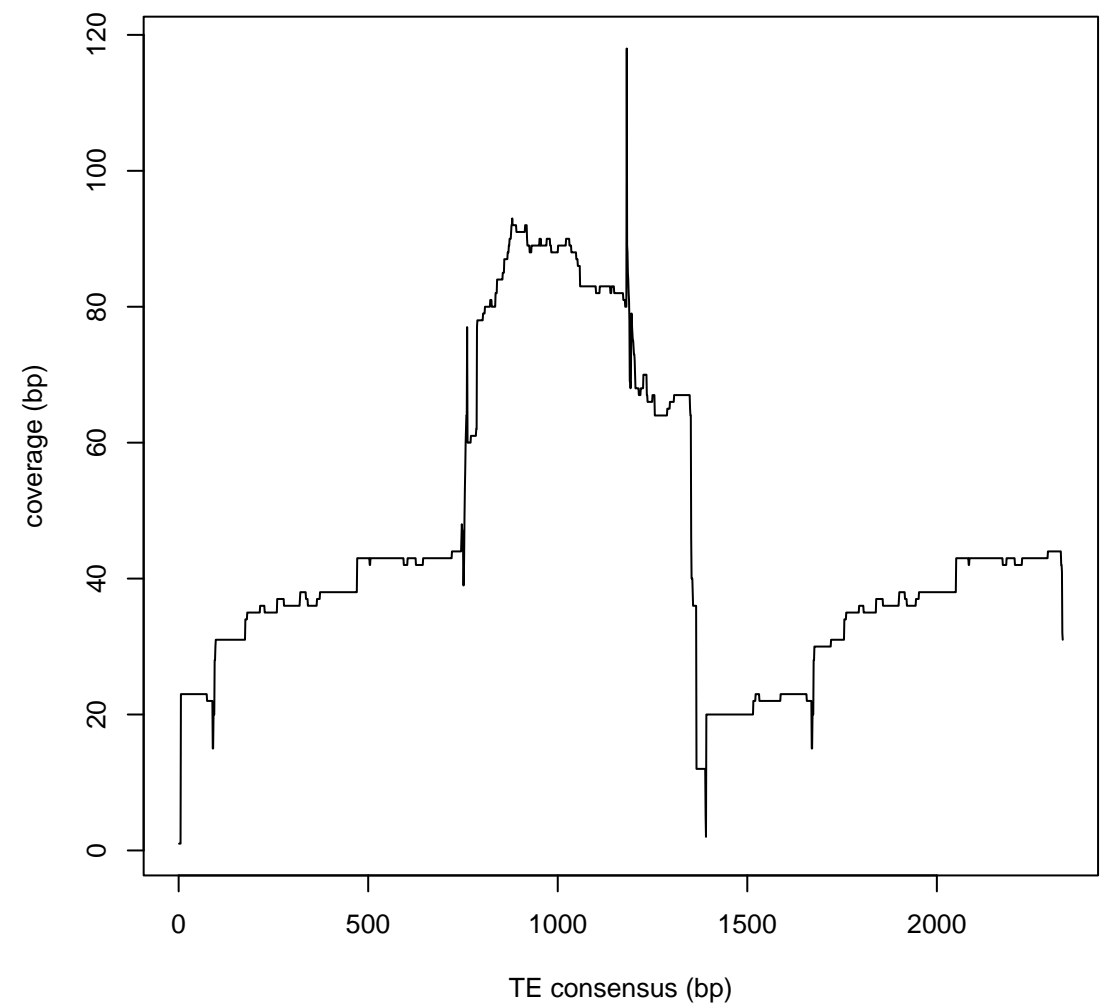


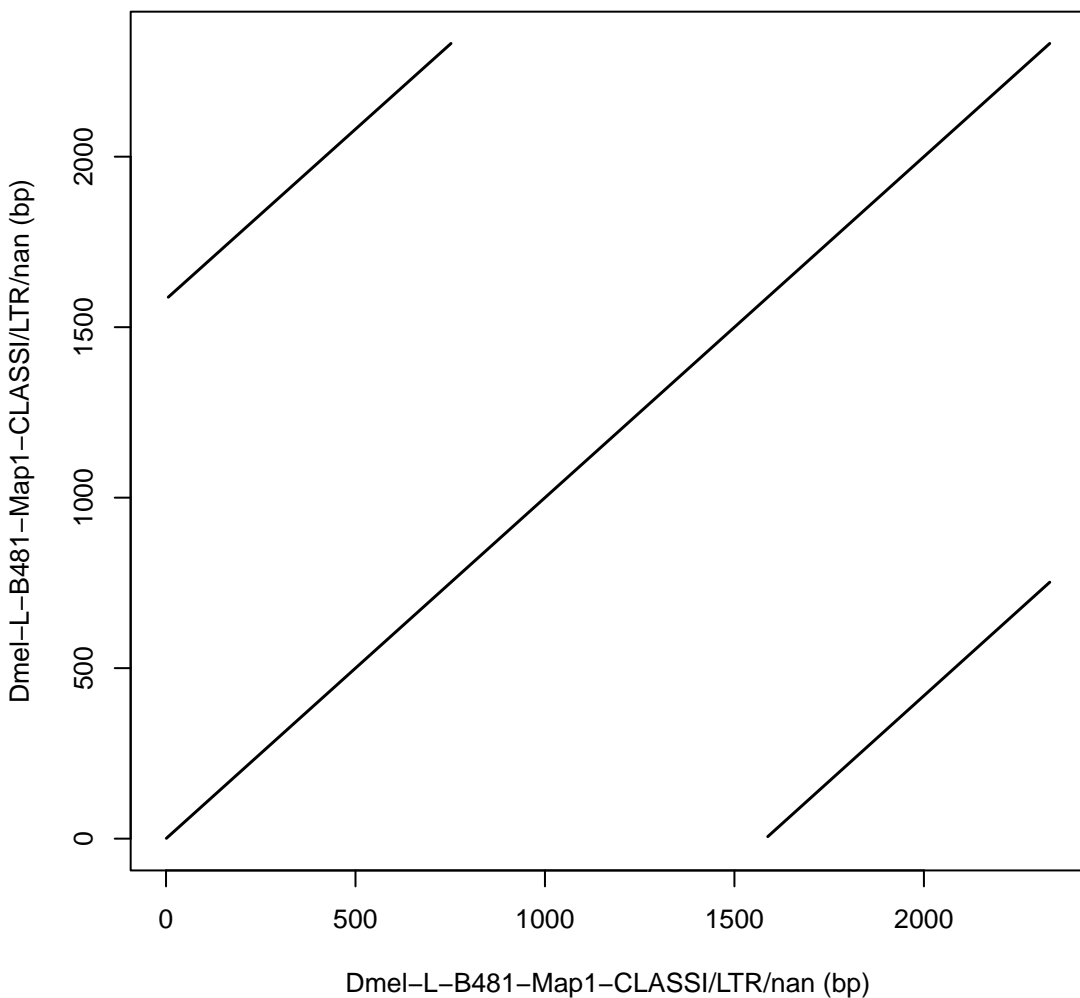
TE: Dmel-L-B481-Map1-CLASSI/LTR/nan
consensus size: 2332bp; fragments: 271; full length: 1 (>=2098.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

