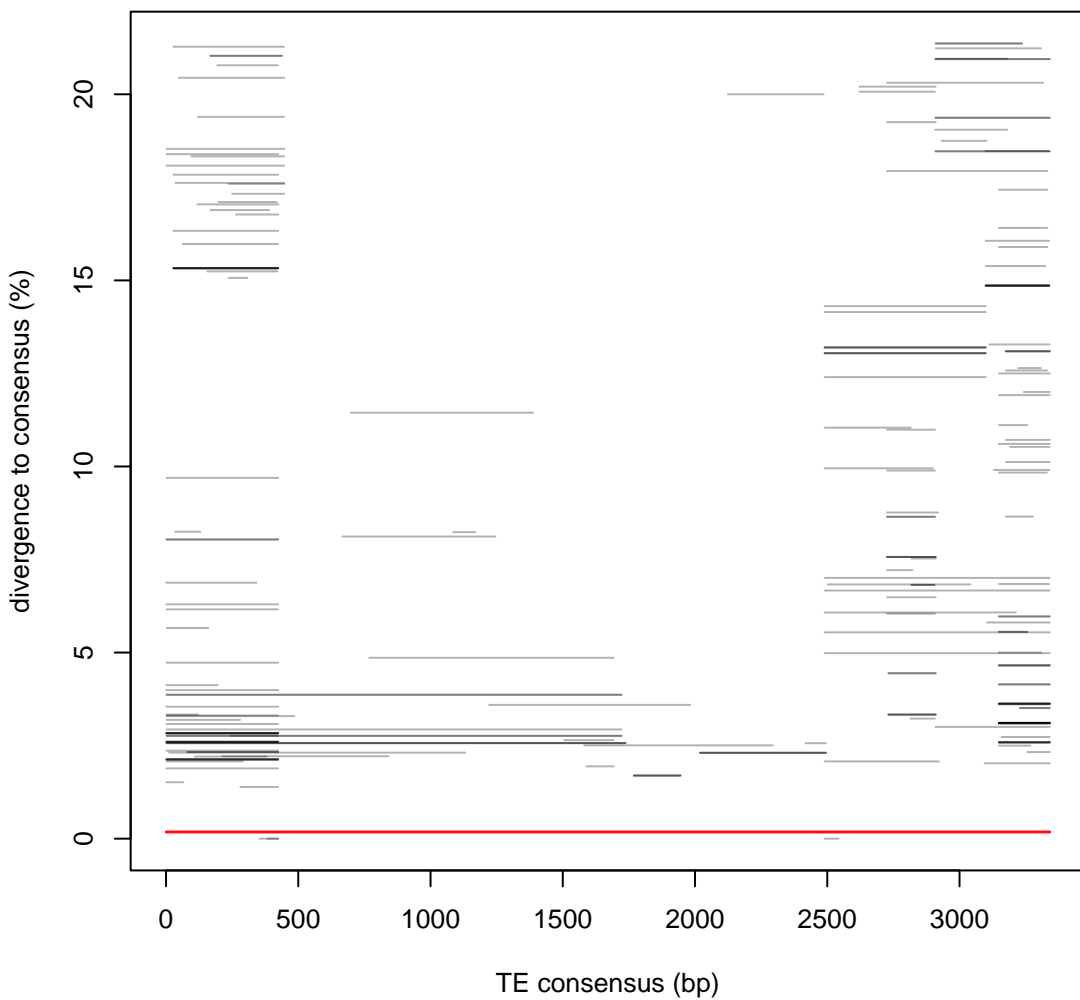
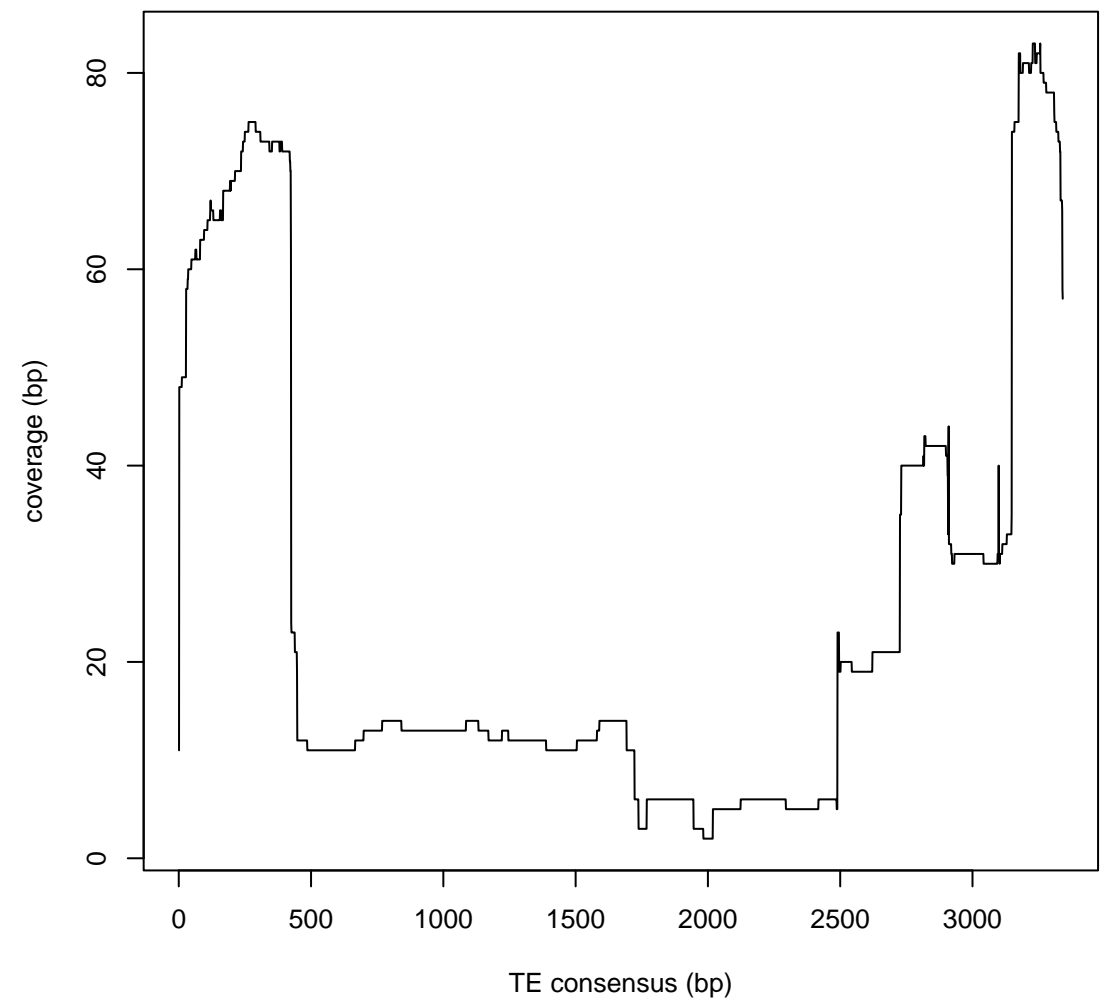


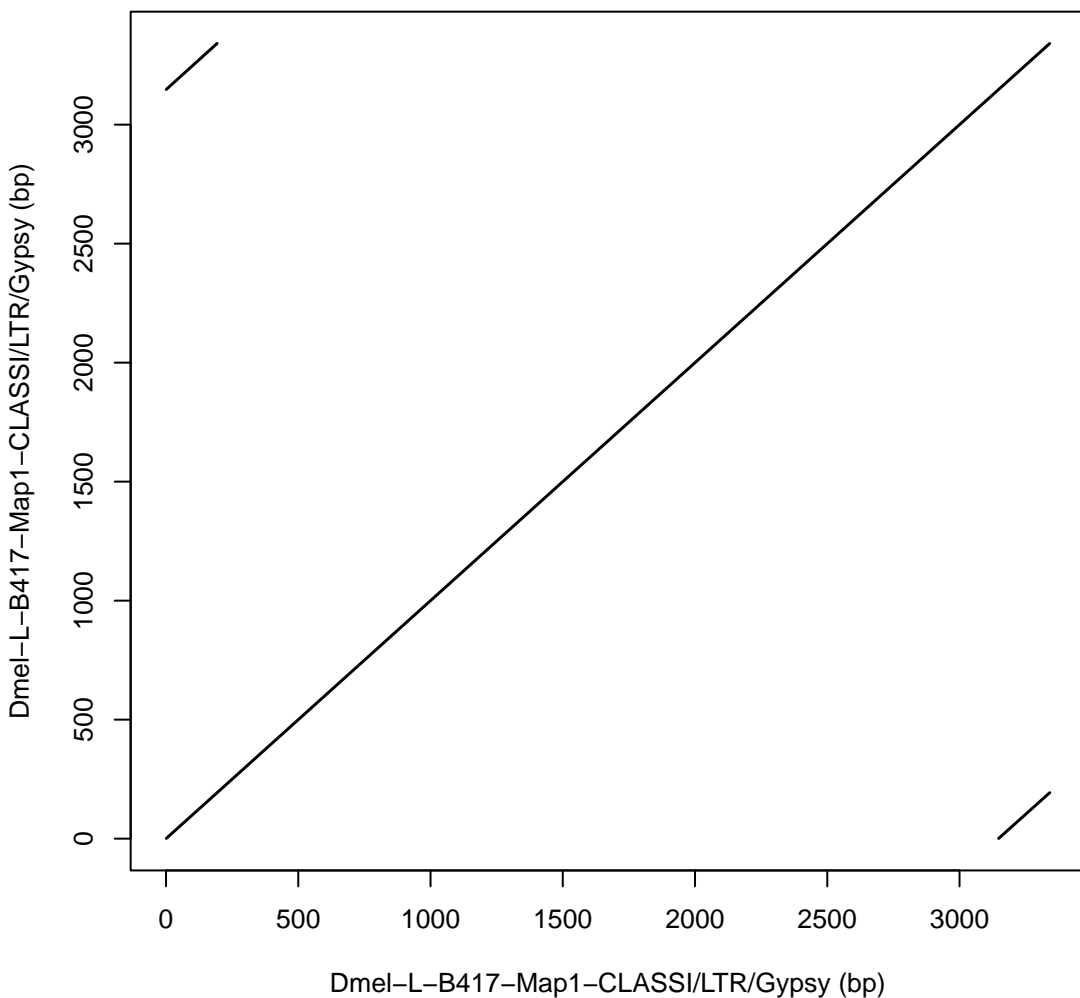
TE: Dmel-L-B417-Map1-CLASSI/LTR/Gypsy
consensus size: 3341bp; fragments: 224; full length: 1 (>=3006.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

