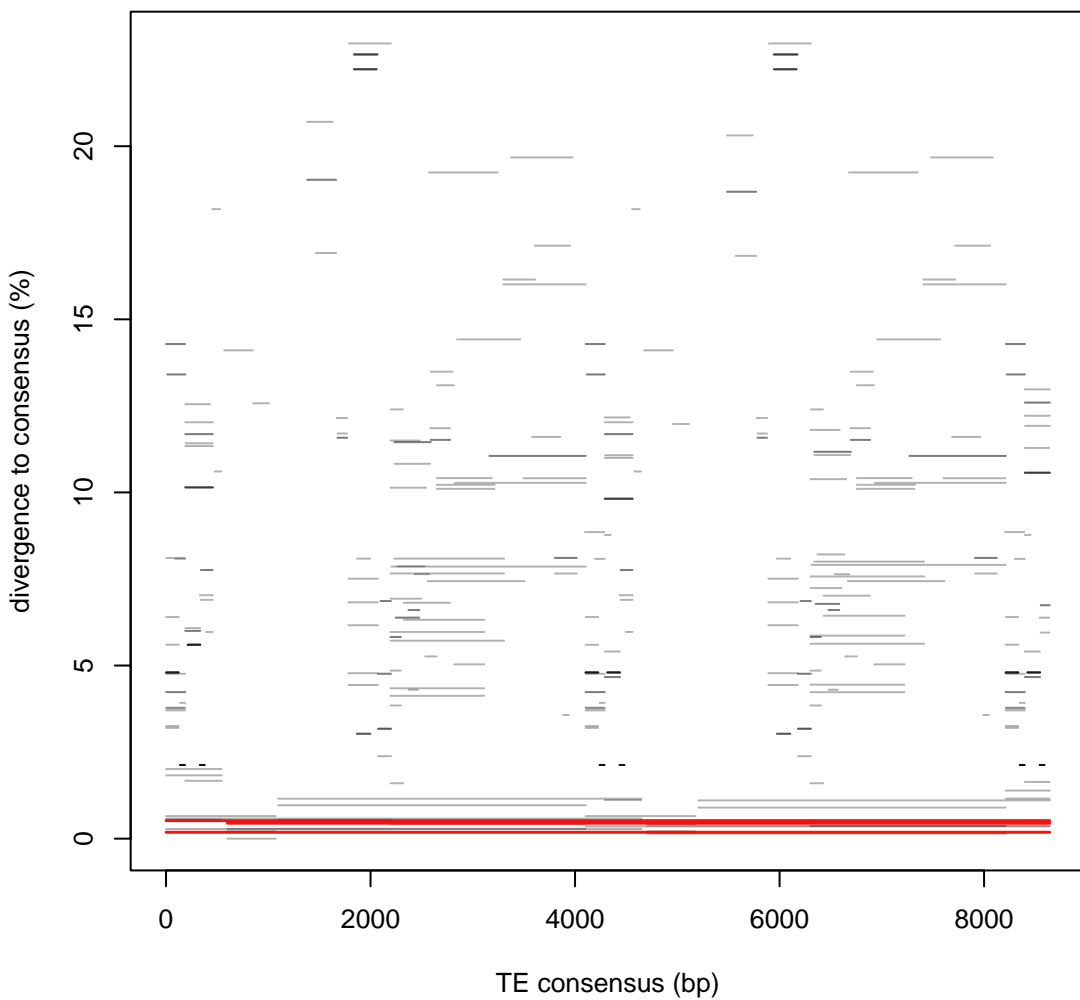
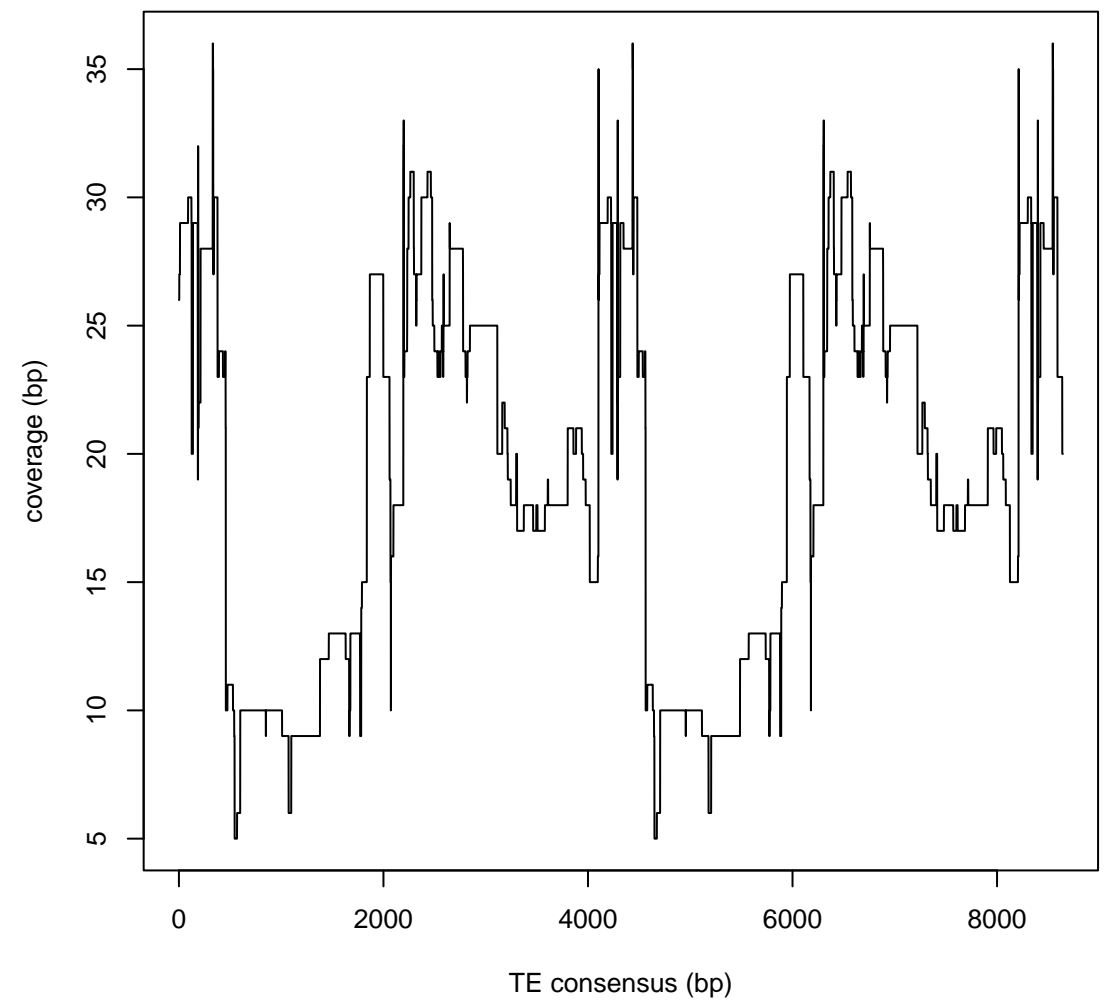


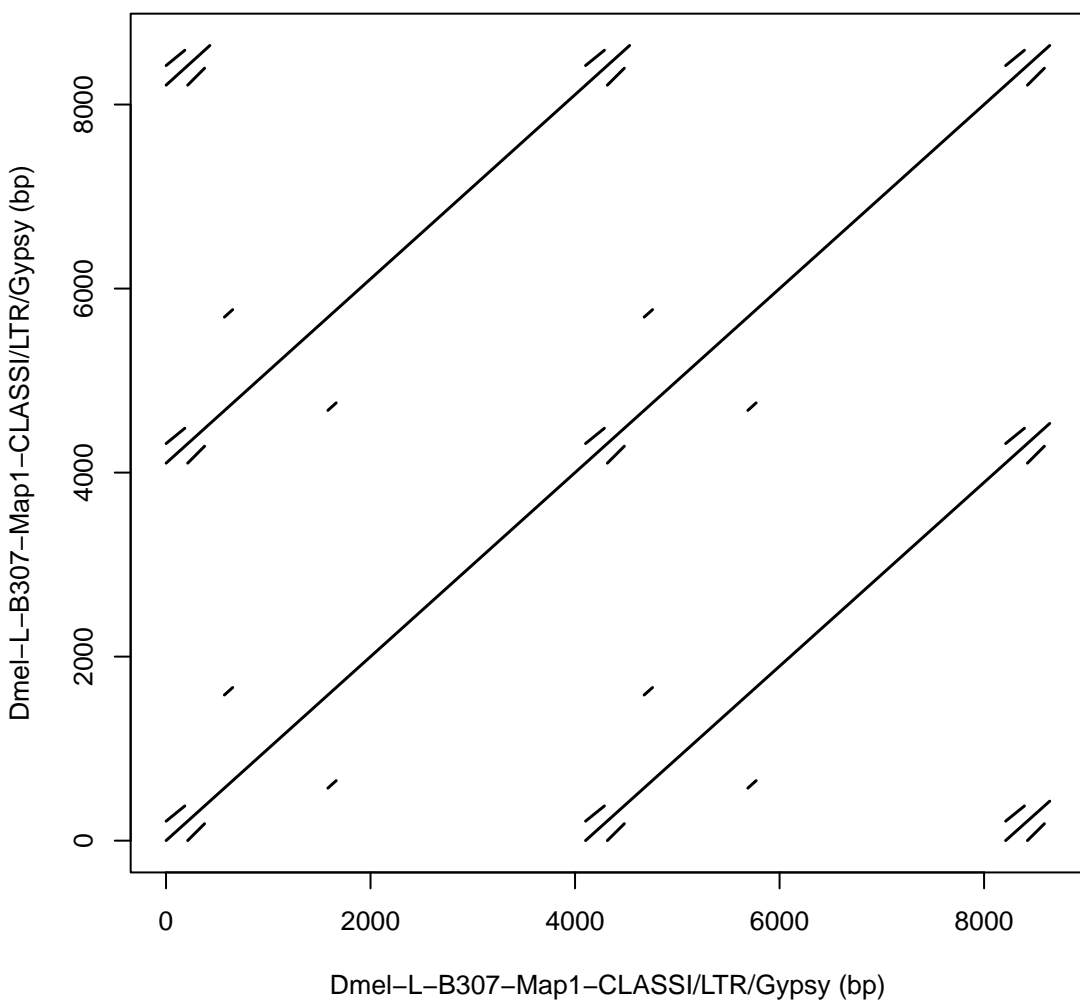
TE: Dmel-L-B307-Map1-CLASSI/LTR/Gypsy
consensus size: 8642bp; fragments: 402; full length: 3 (>=7777.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

