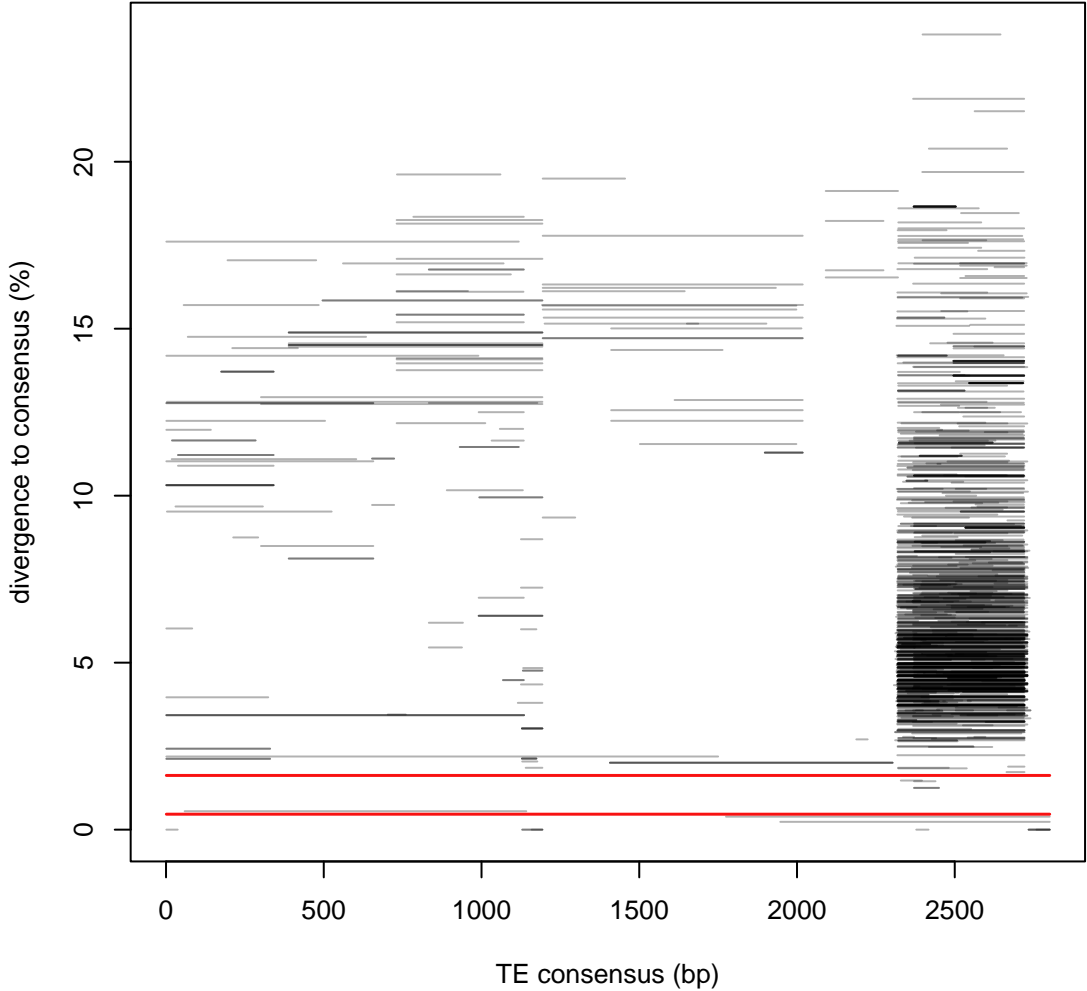
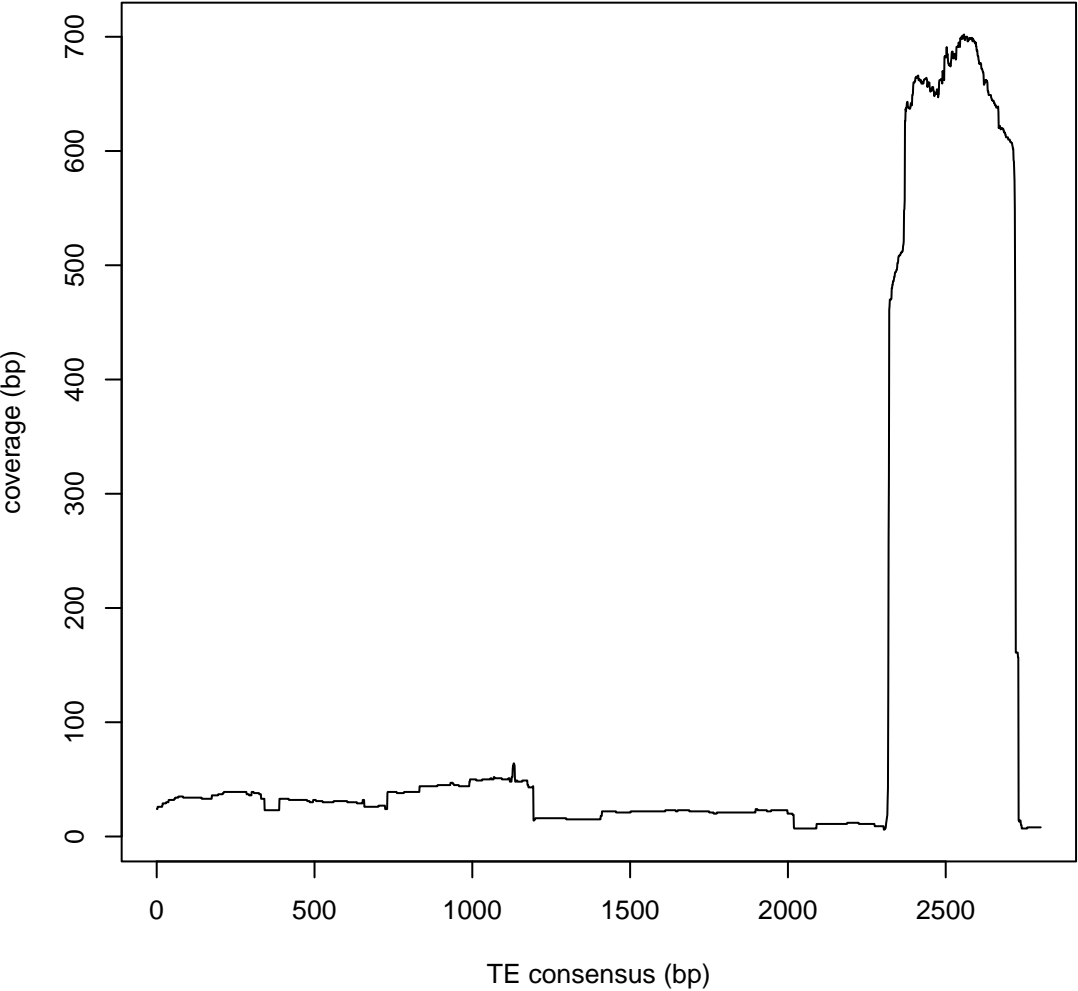


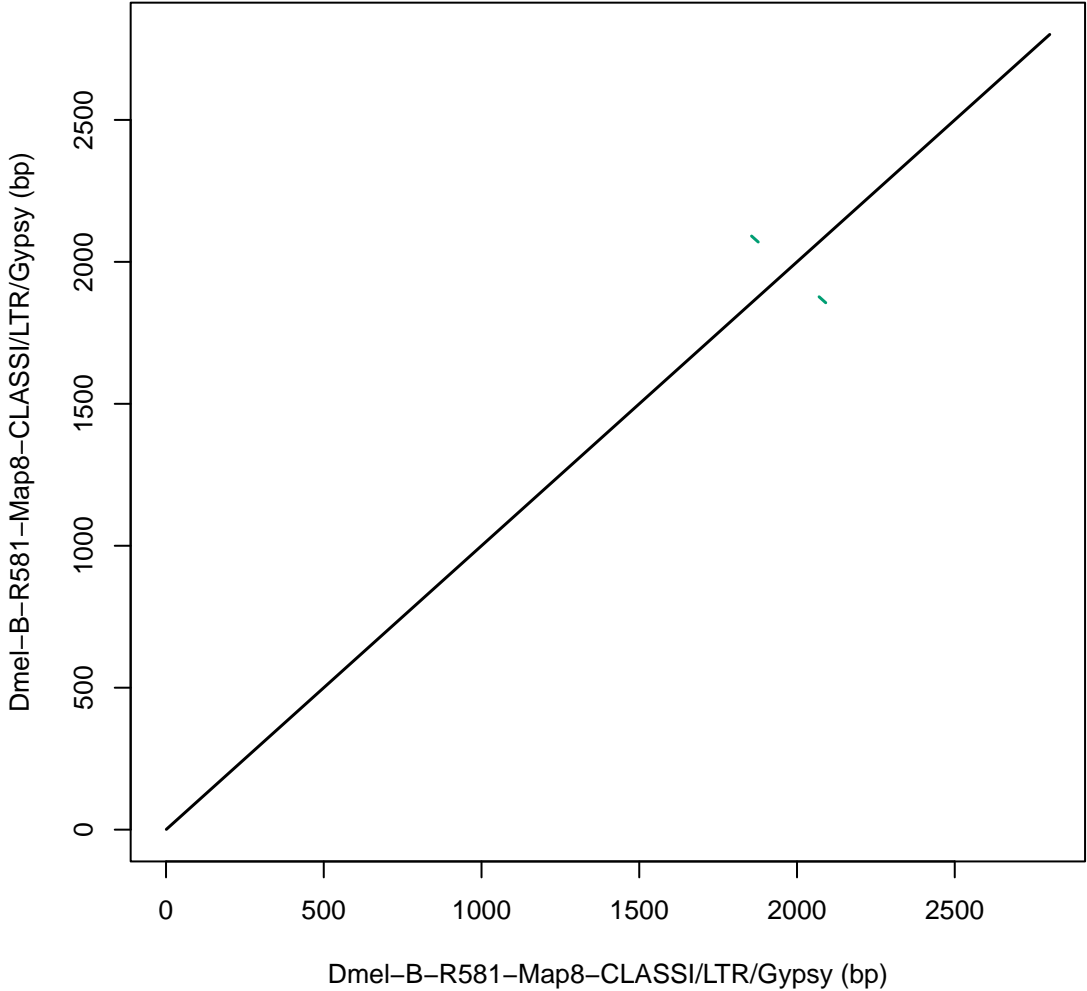
TE: Dmel-B-R581-Map8-CLASSI/LTR/Gypsy
consensus size: 2801bp; fragments: 1031; full length: 2 (>=2520.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

