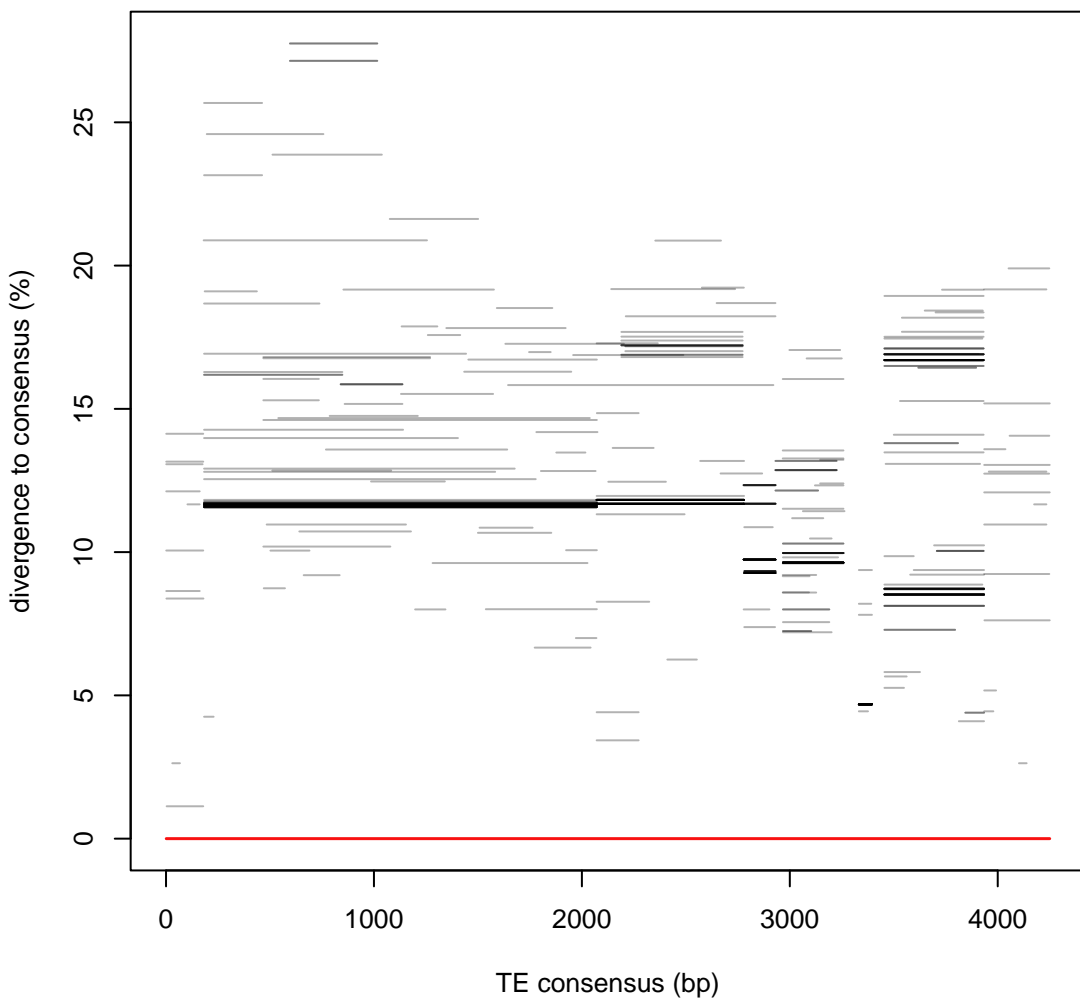
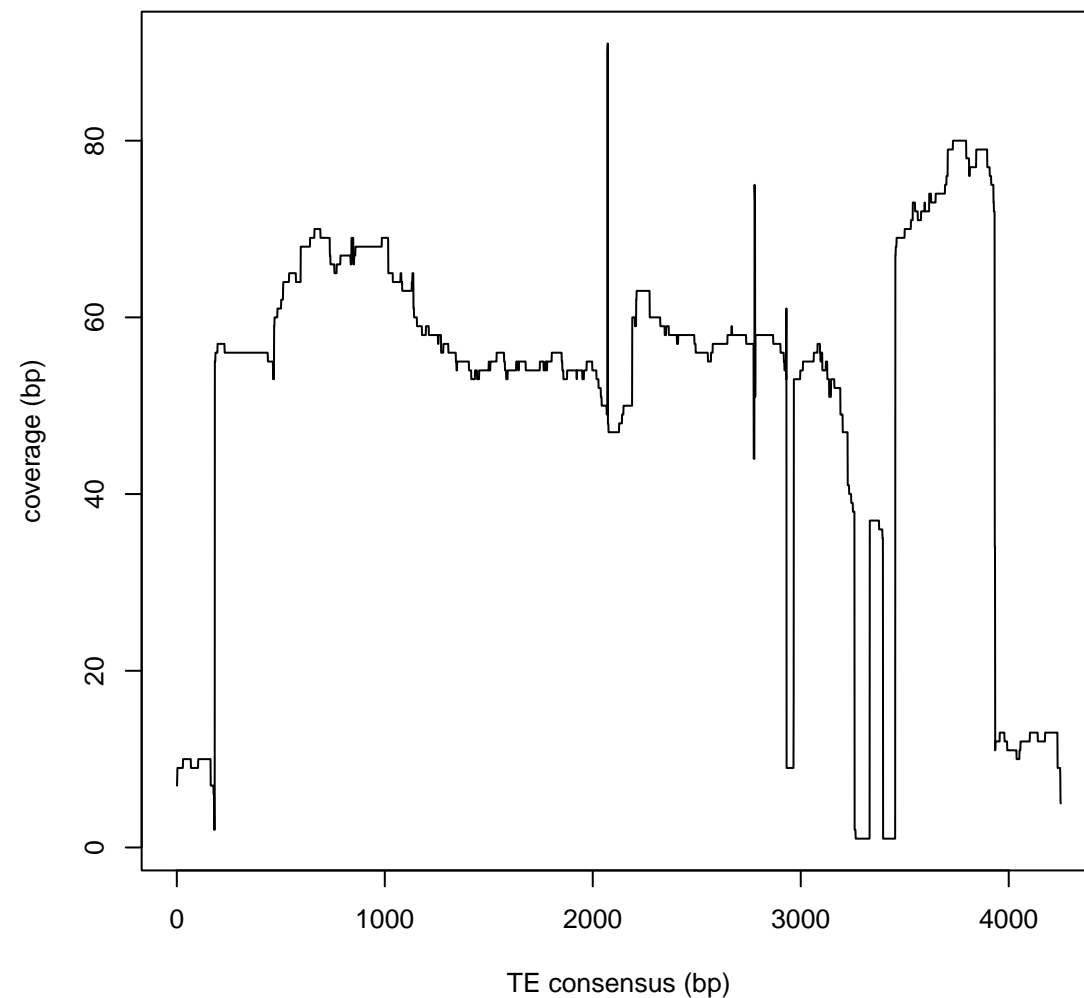


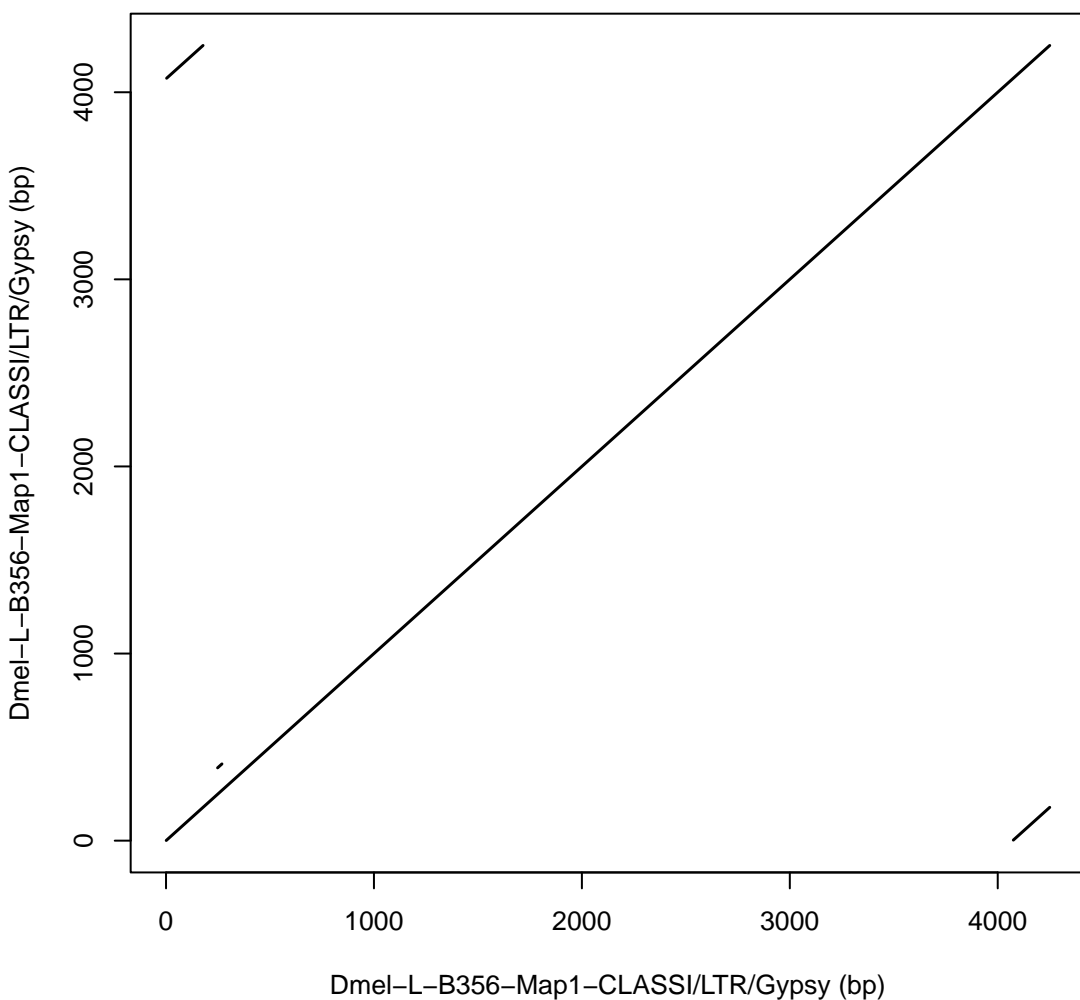
TE: Dmel-L-B356-Map1-CLASSI/LTR/Gypsy
consensus size: 4250bp; fragments: 433; full length: 1 (>=3825bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

