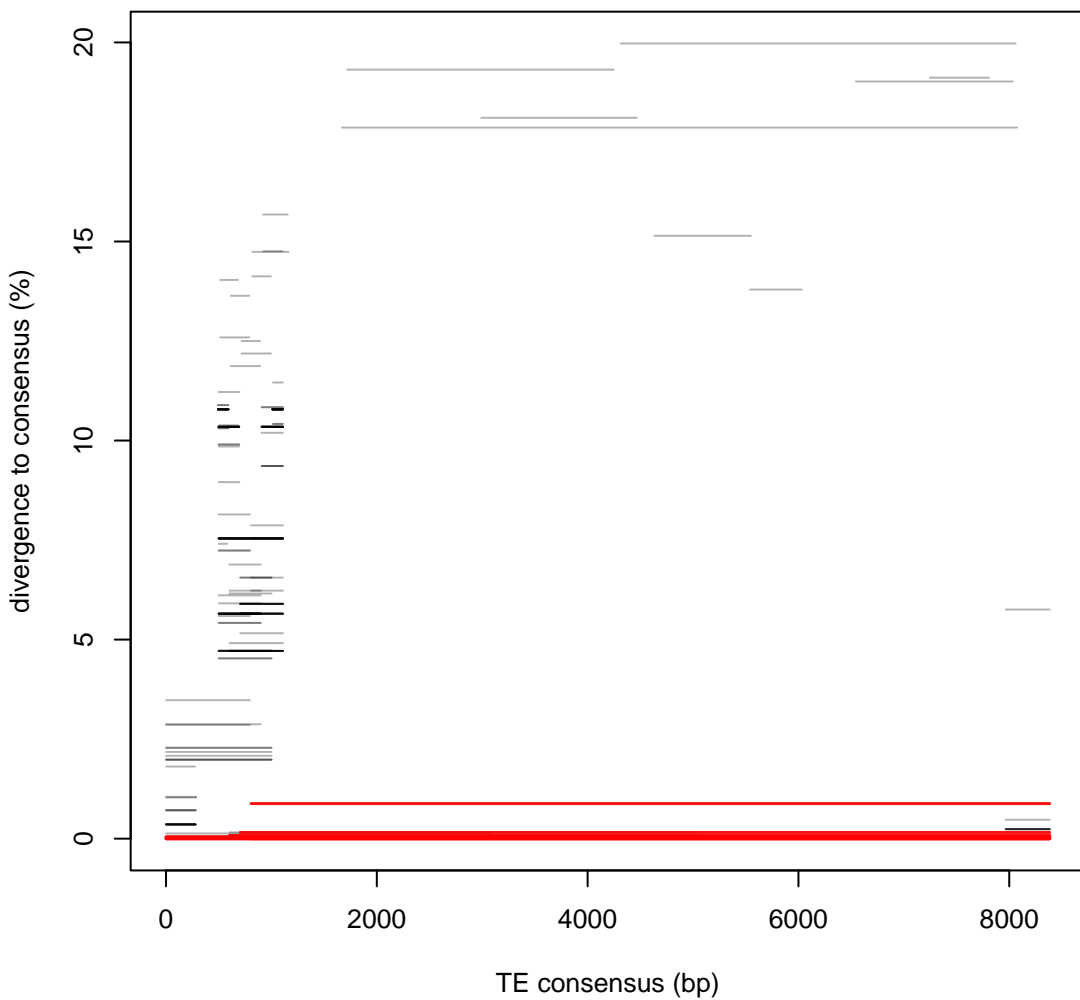
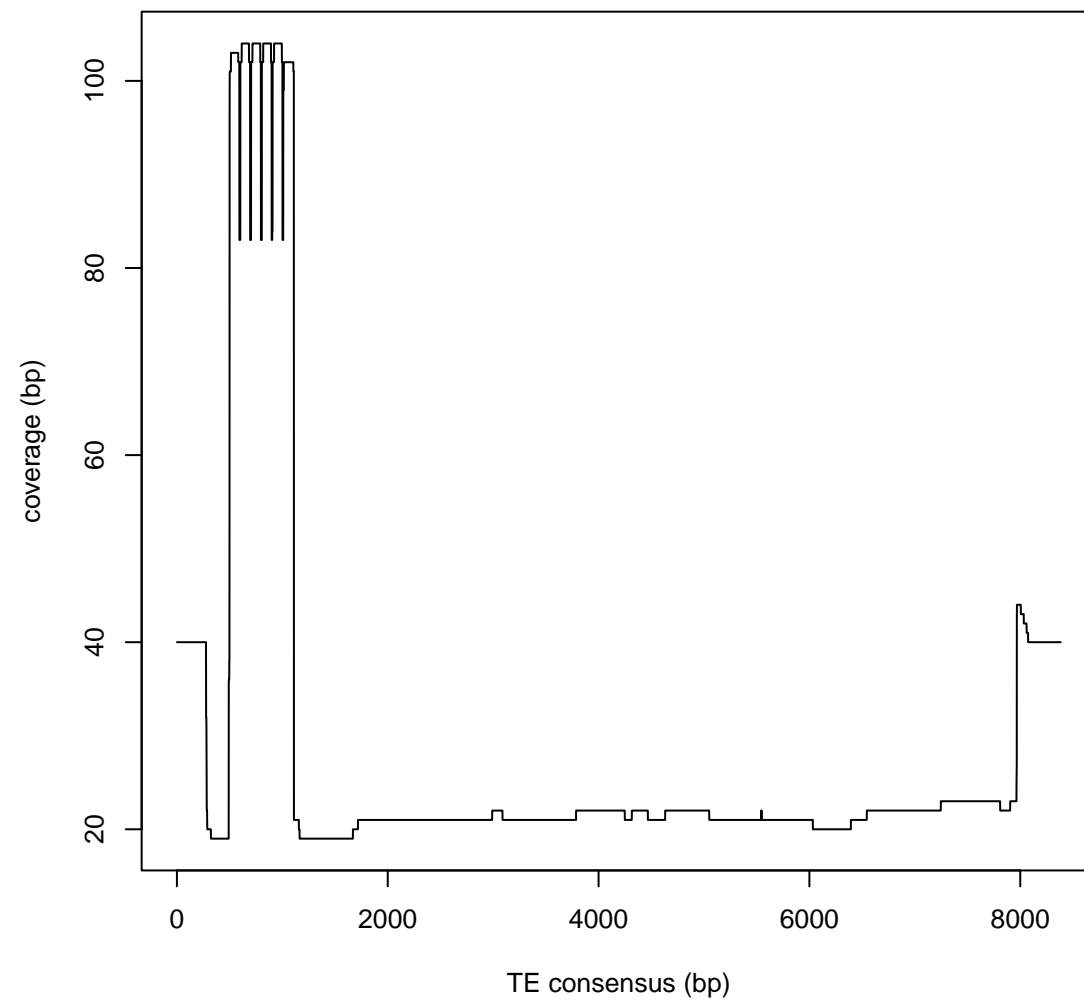


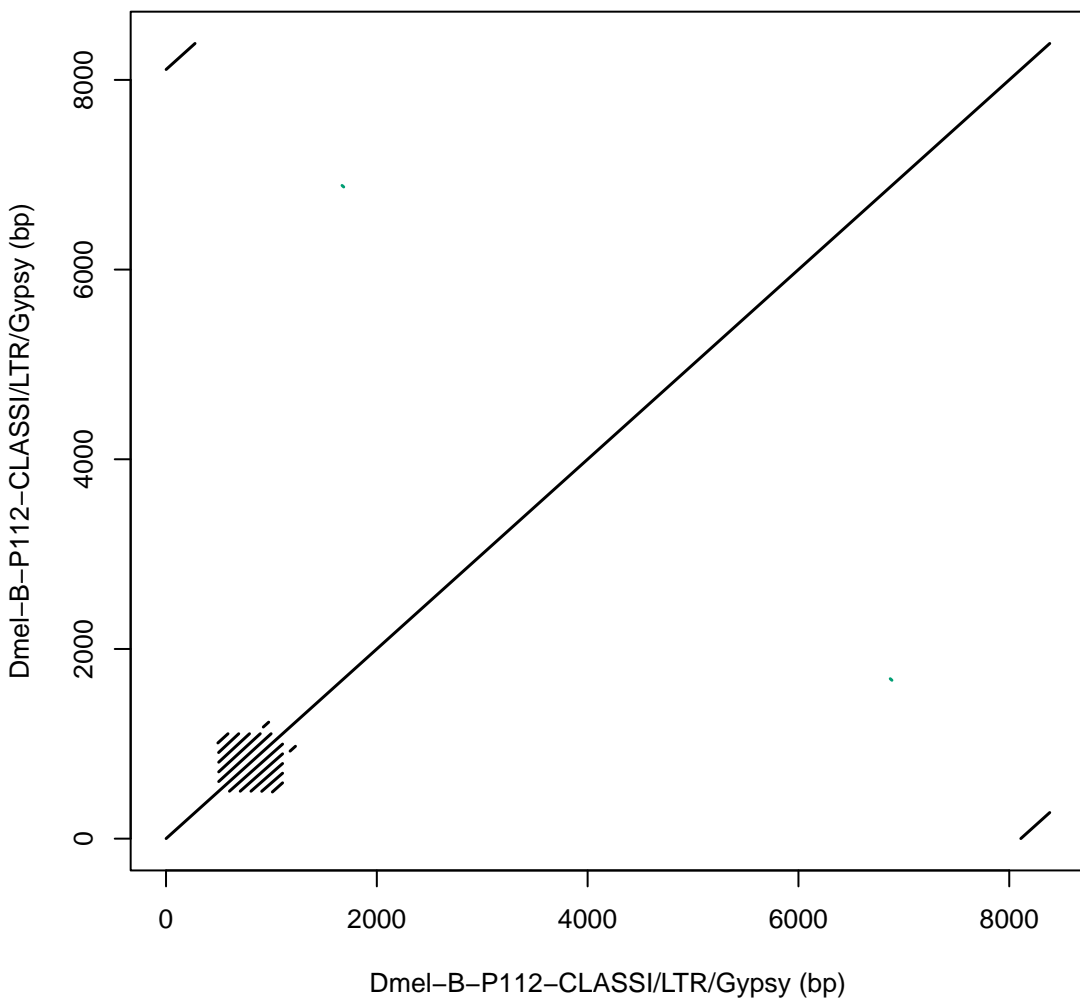
TE: Dmel-B-P112-CLASSI/LTR/Gypsy
consensus size: 8384bp; fragments: 259; full length: 17 (>=7545.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

