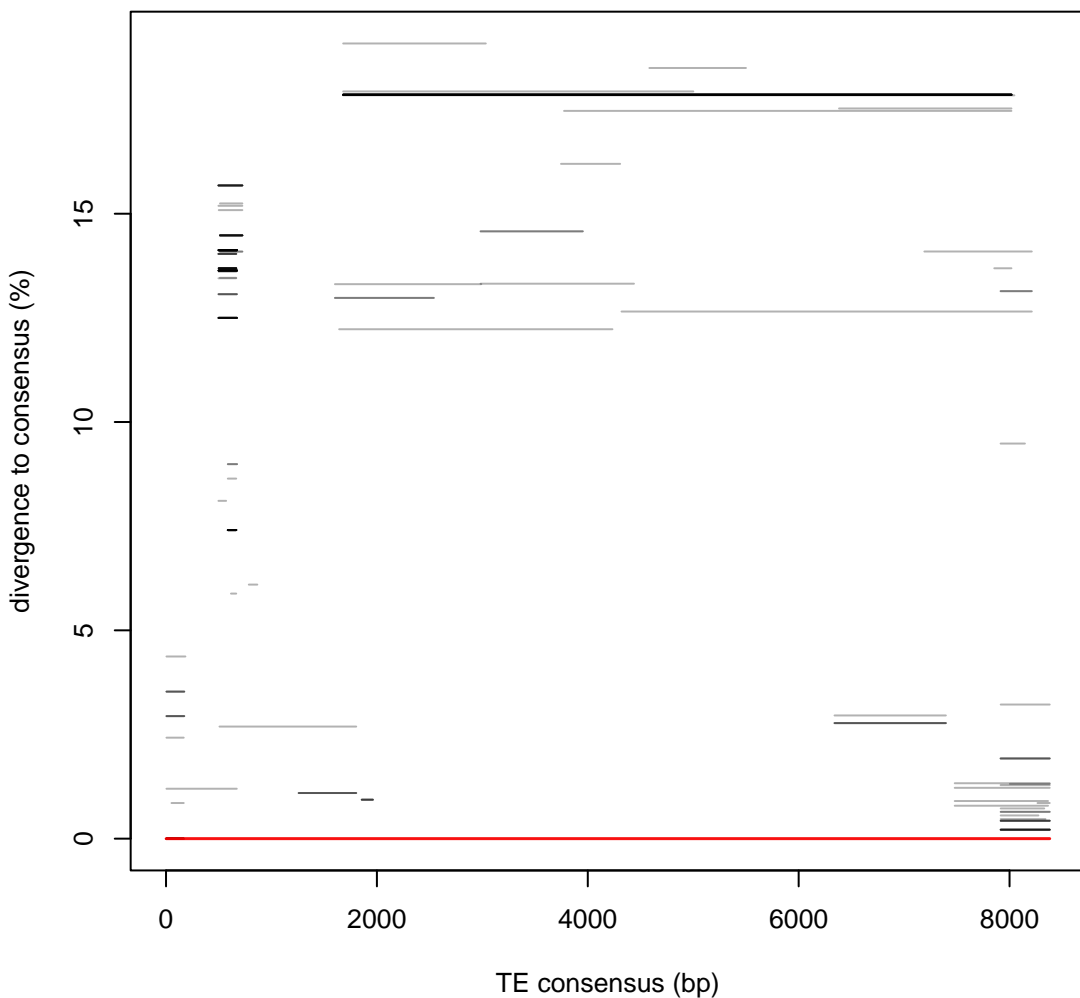
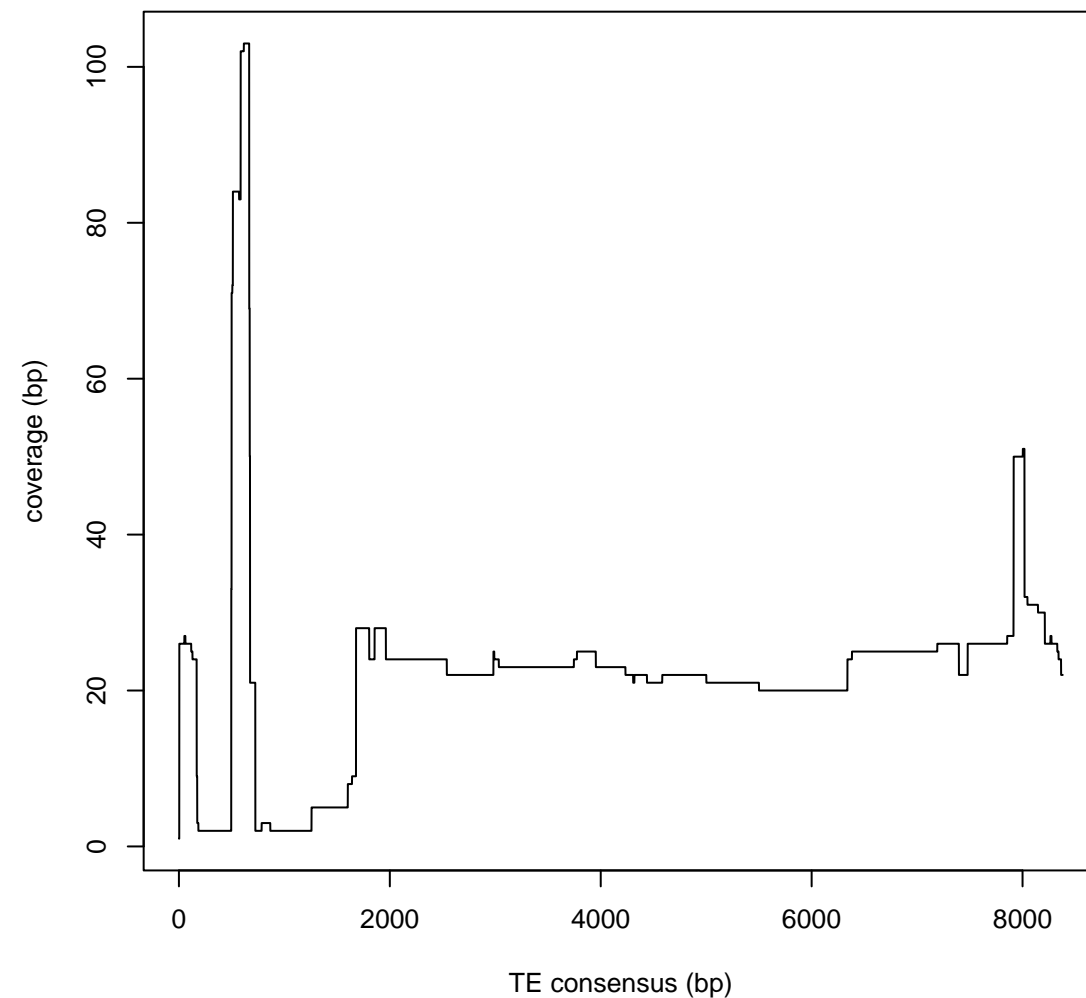


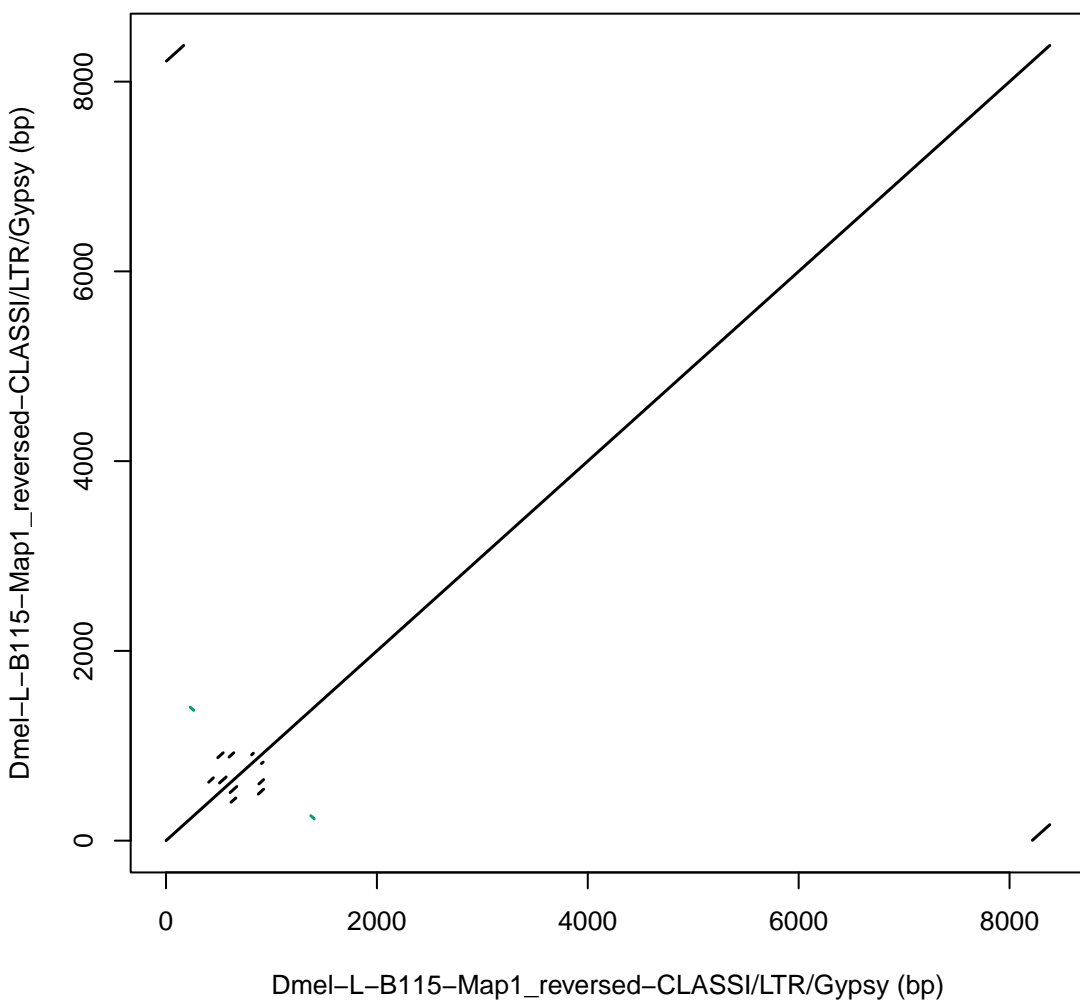
TE: Dmel-L-B115-Map1_reversed-CLASSI/LTR/Gypsy
consensus size: 8381bp; fragments: 203; full length: 1 (>=7542.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

