TE: Dmel−B−R834−Map6−CLASSI/LTR/Gypsy
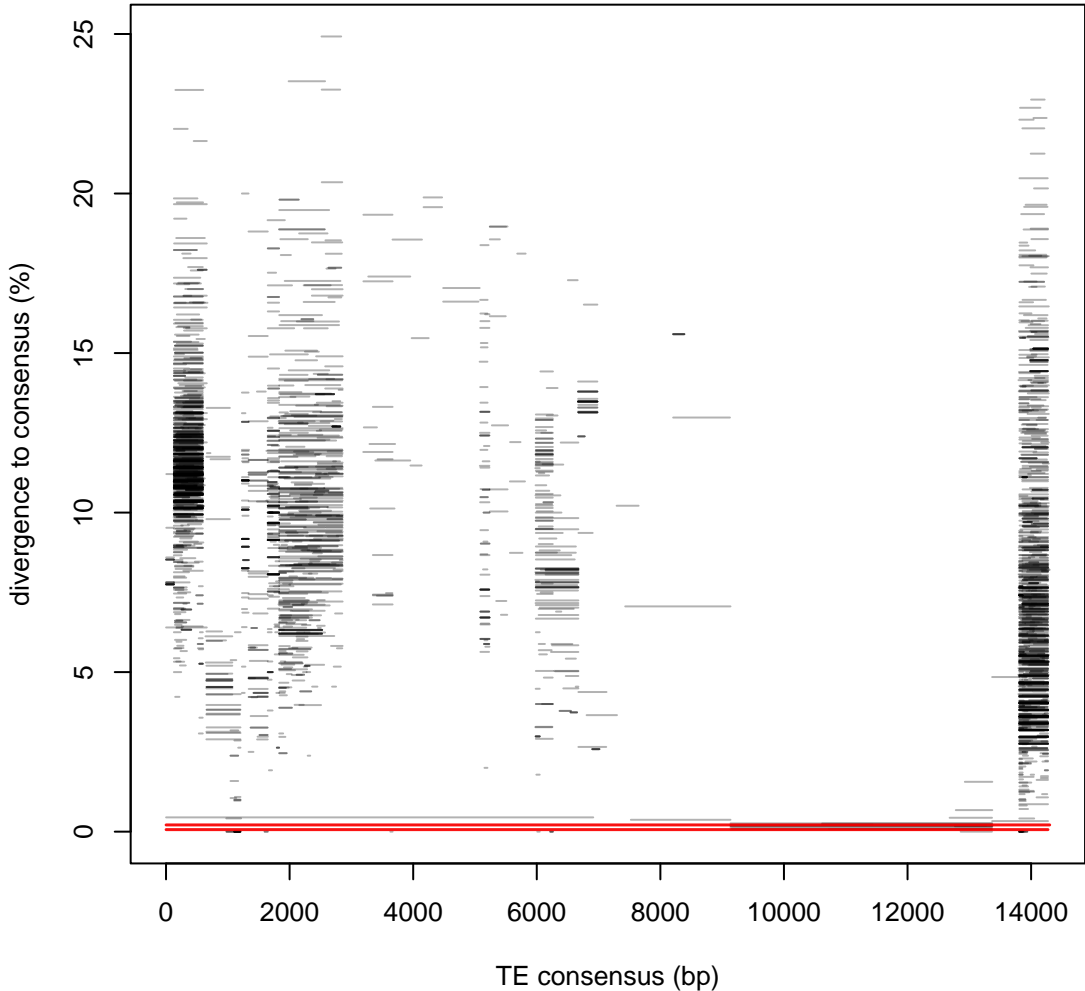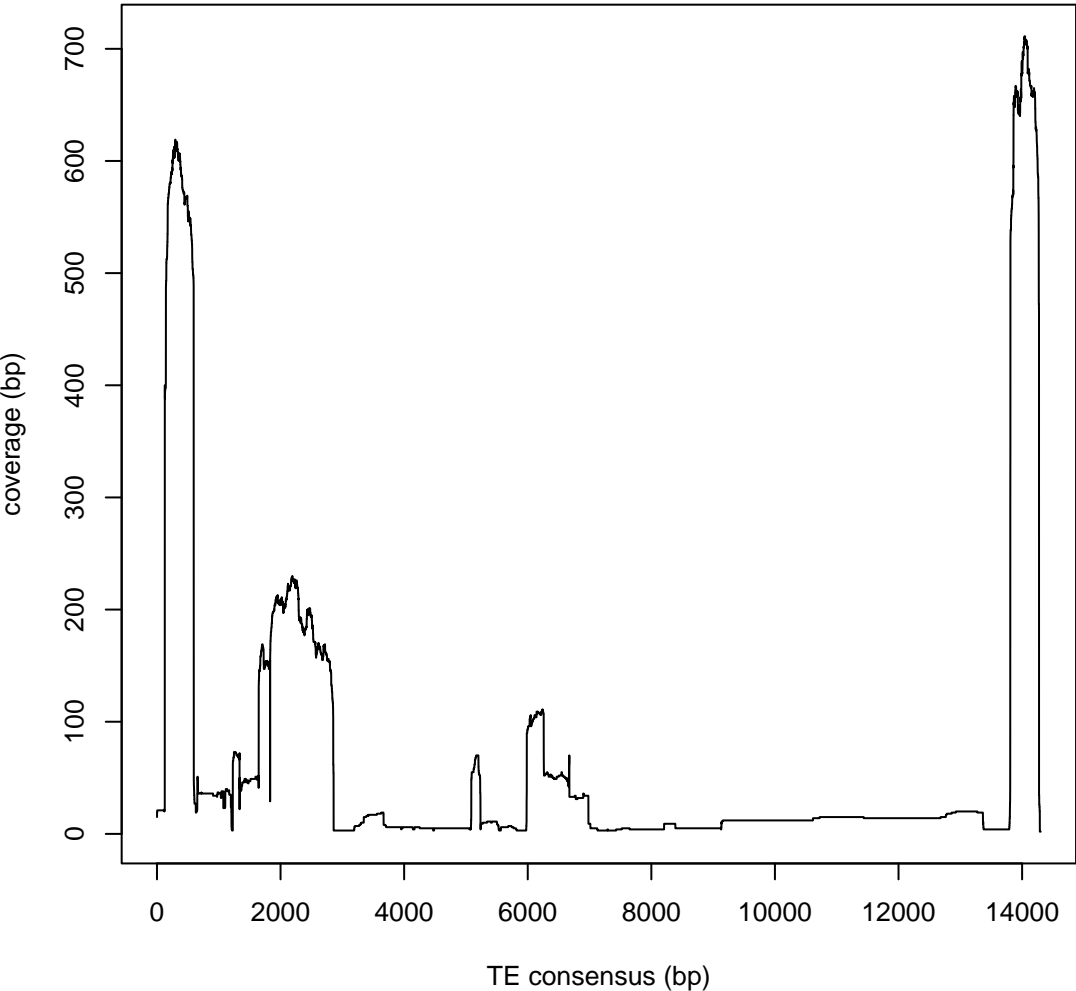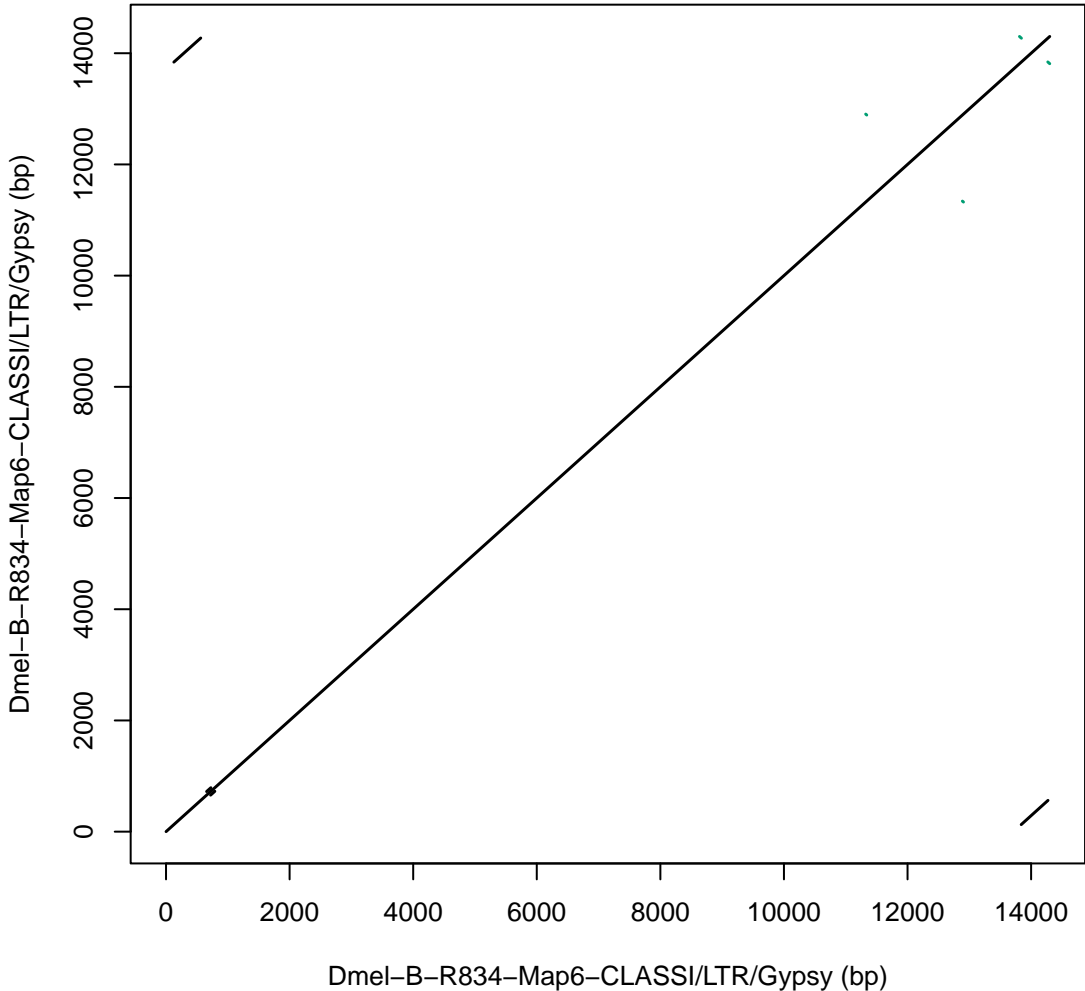consensus size: 14301bp; fragments: 2822; full length: 2 (>=12870.9bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected