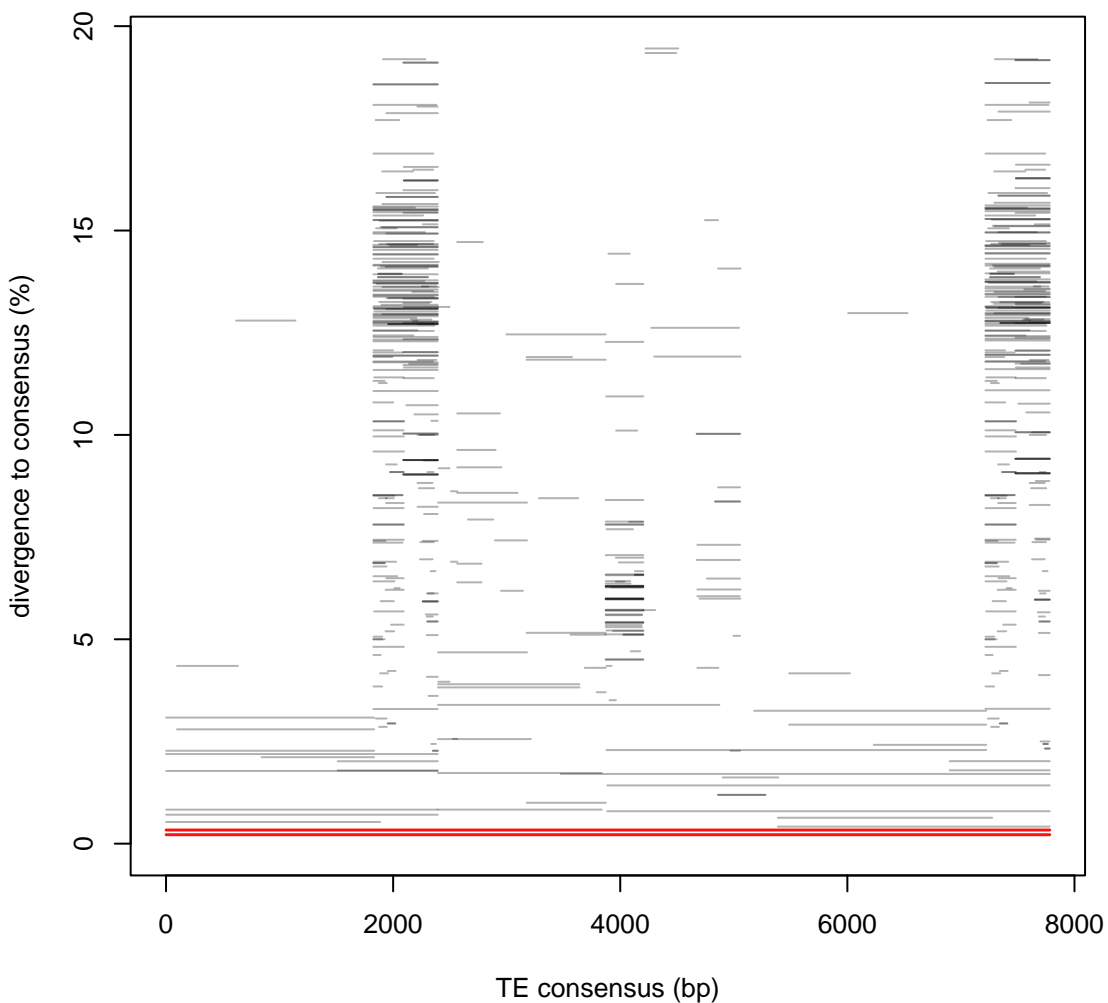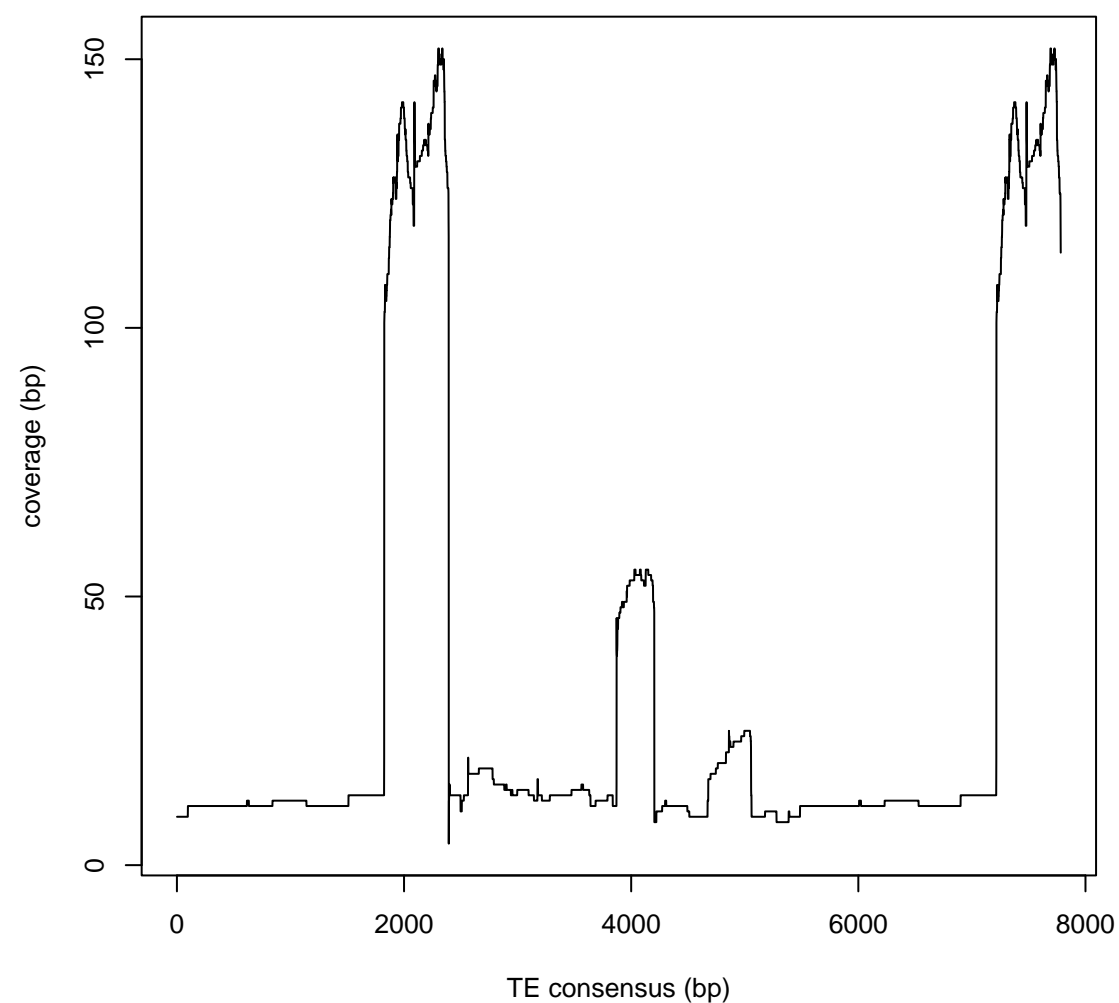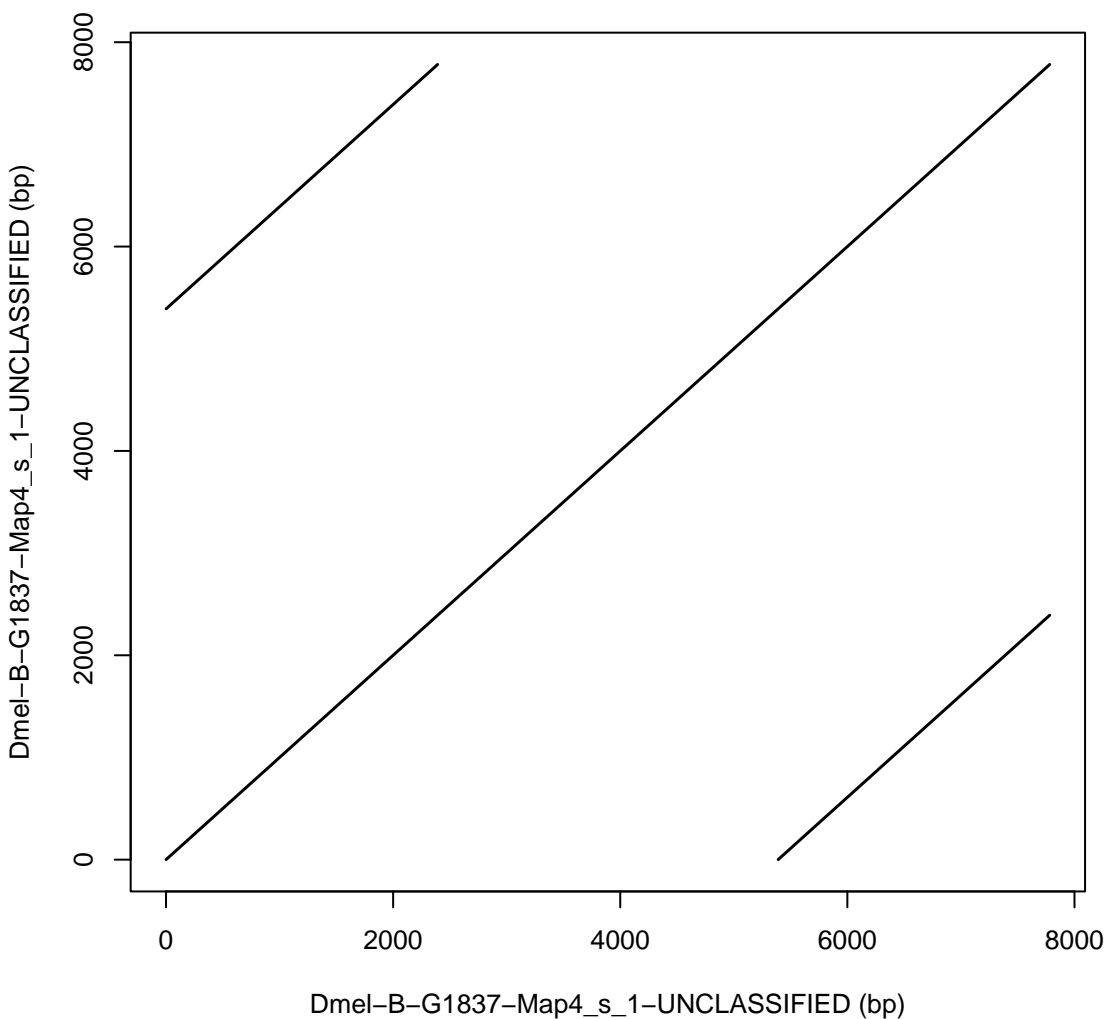## TE: Dmel−B−G1837−Map4_s_1−UNCLASSIFIED
consensus size: 7782bp; fragments: 578; full length: 2 (>=7003.8bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

CR1−1_DRh_gag LINE/CR1
Transib−10_HM TIR/CMC−Transib
CR1−1_DRh_gag LI