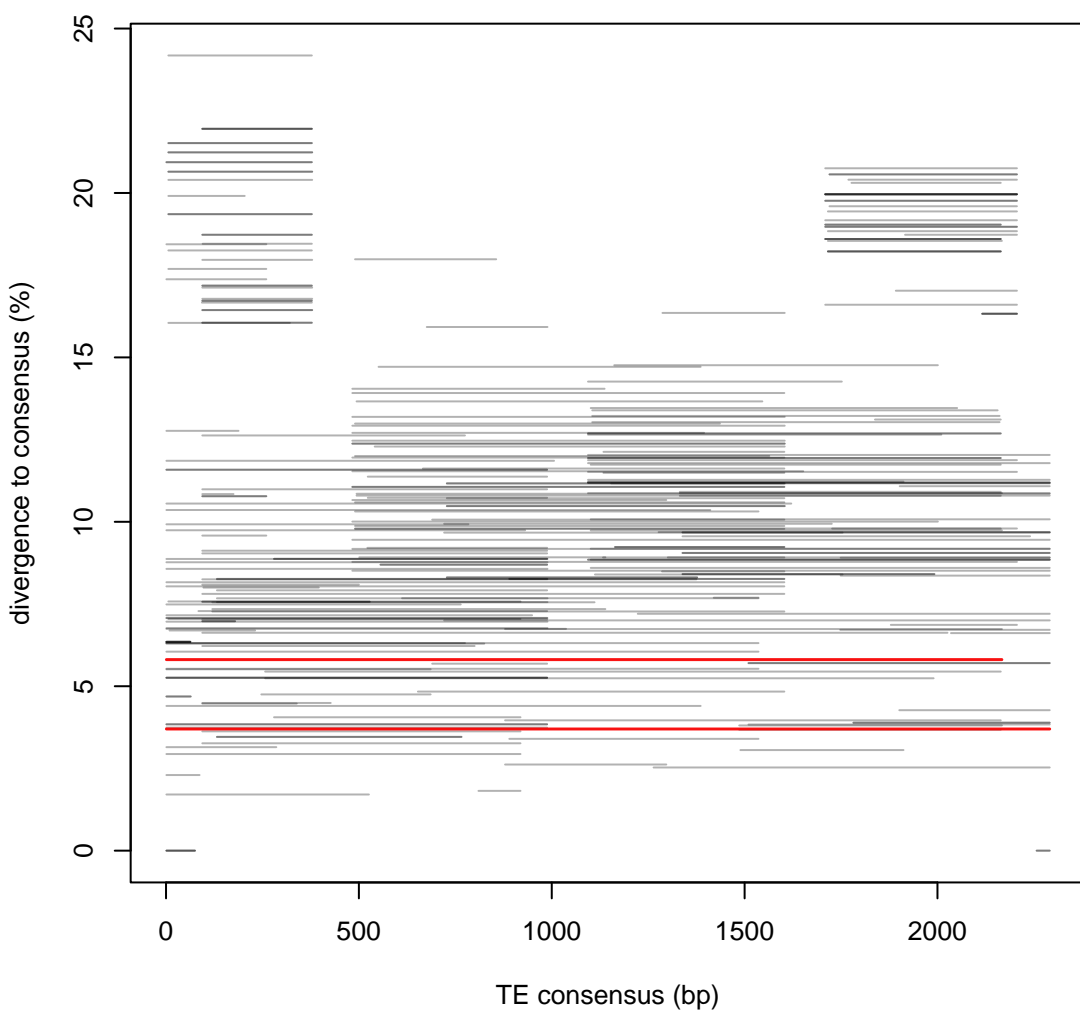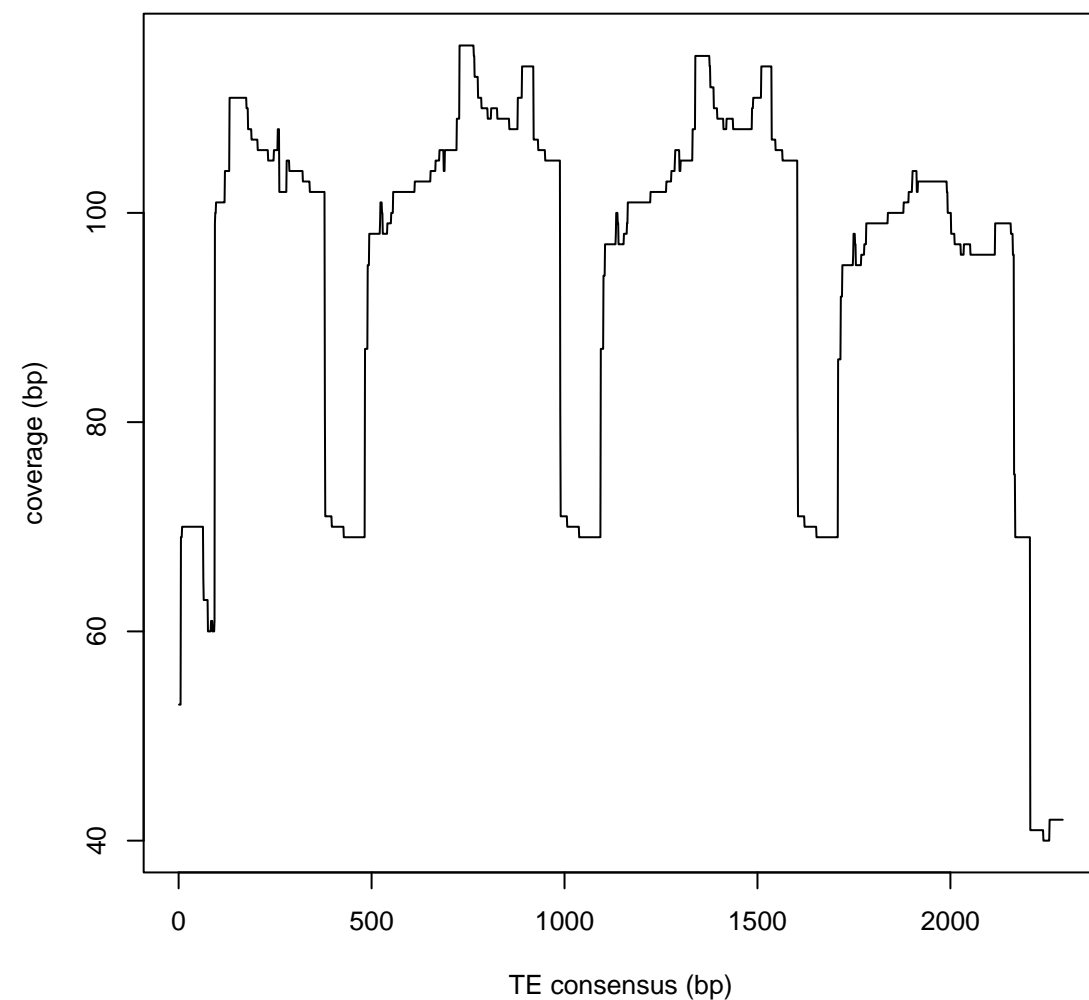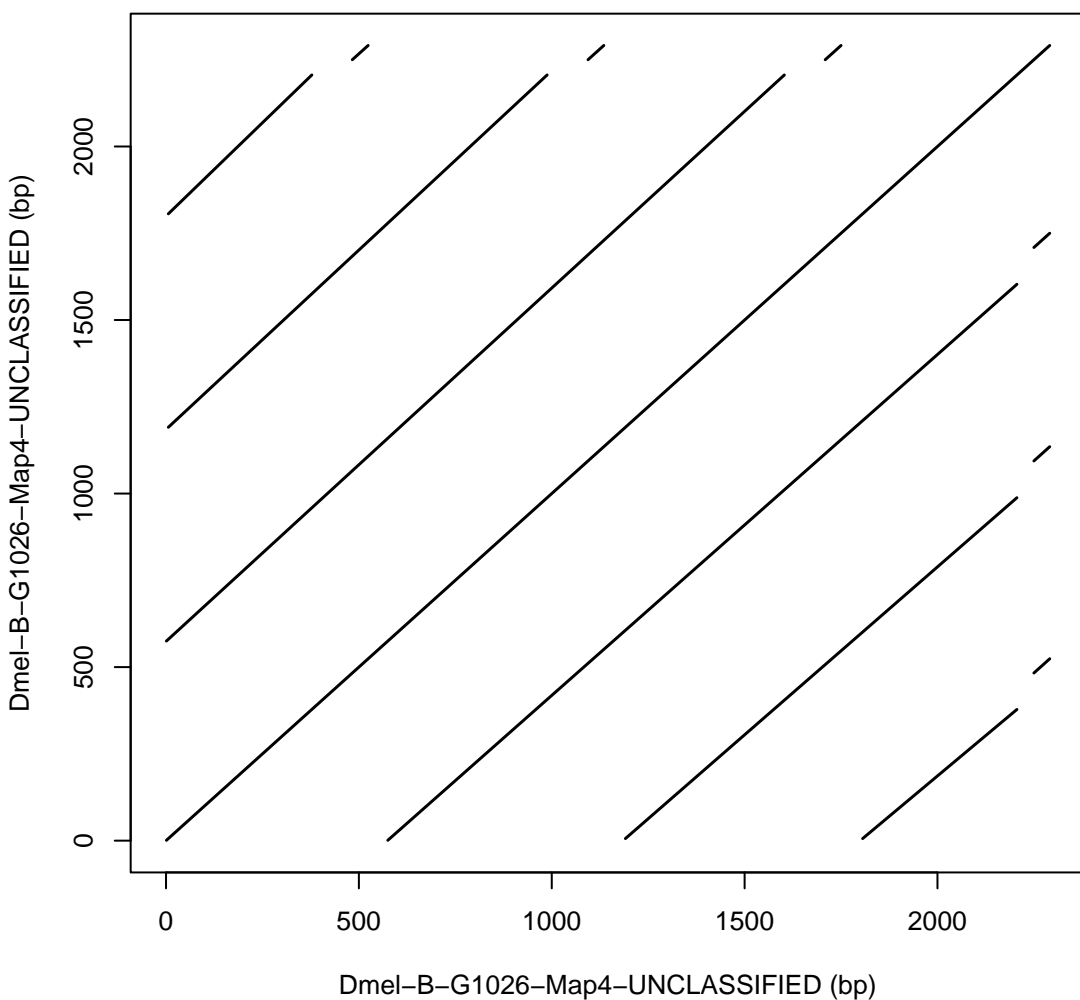**TE: Dmel−B−G1026−Map4−UNCLASSIFIED**
consensus size: 2291bp; fragments: 293; full length: 2 (>=2061.9bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected