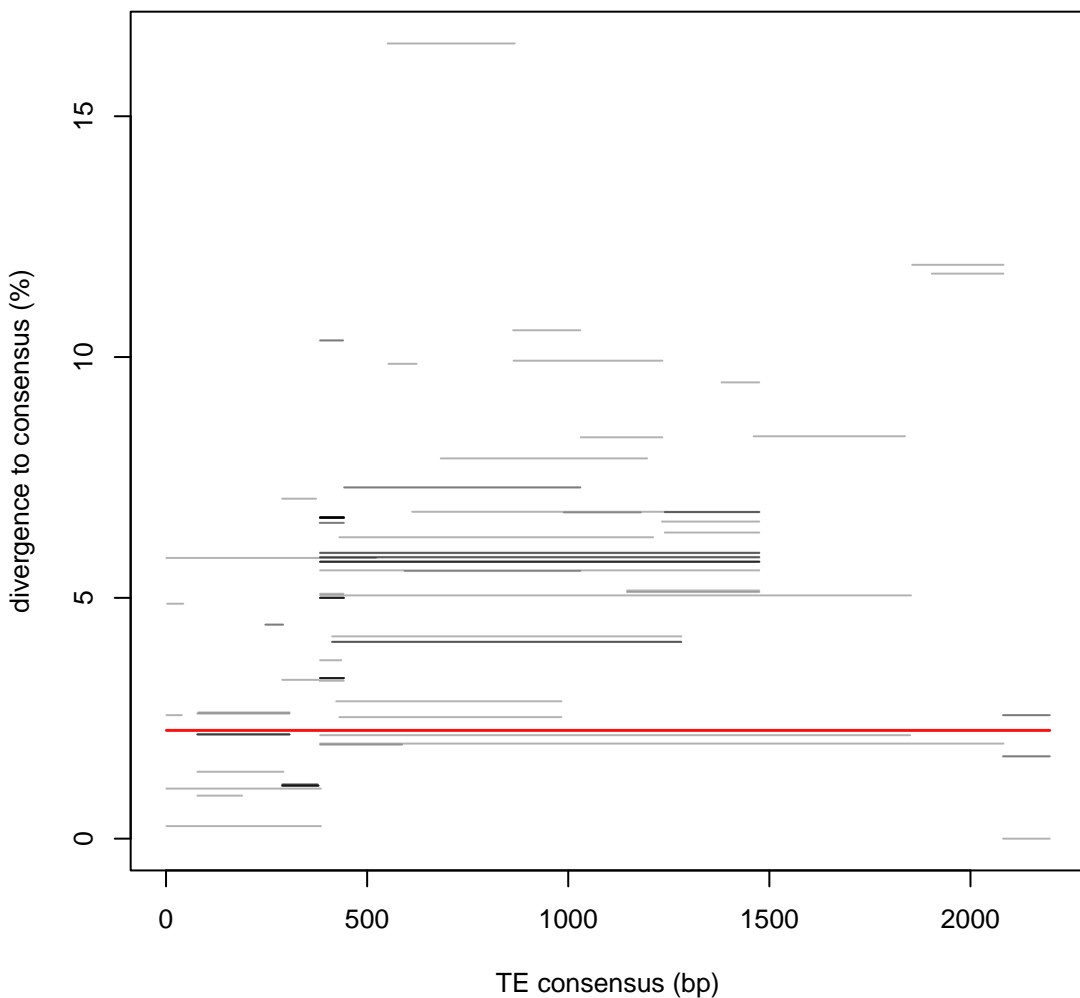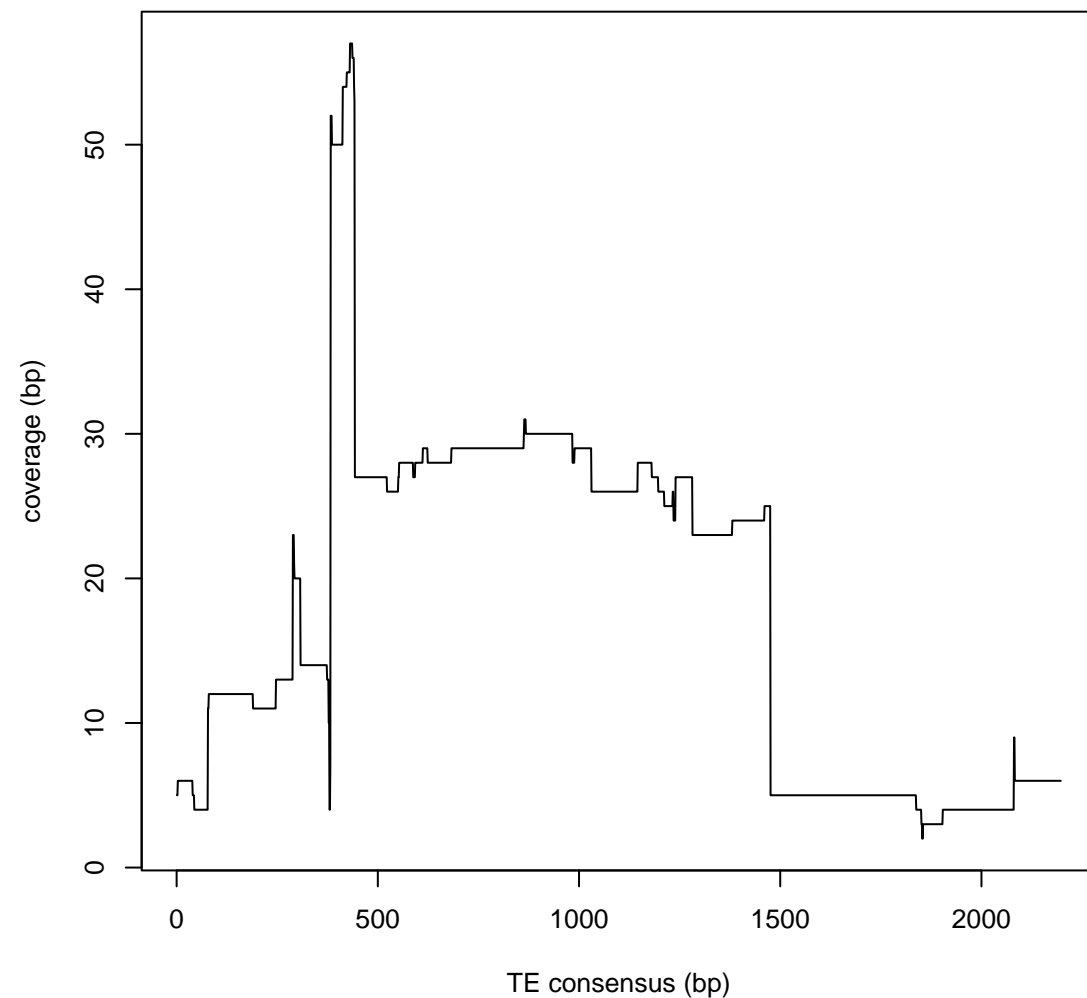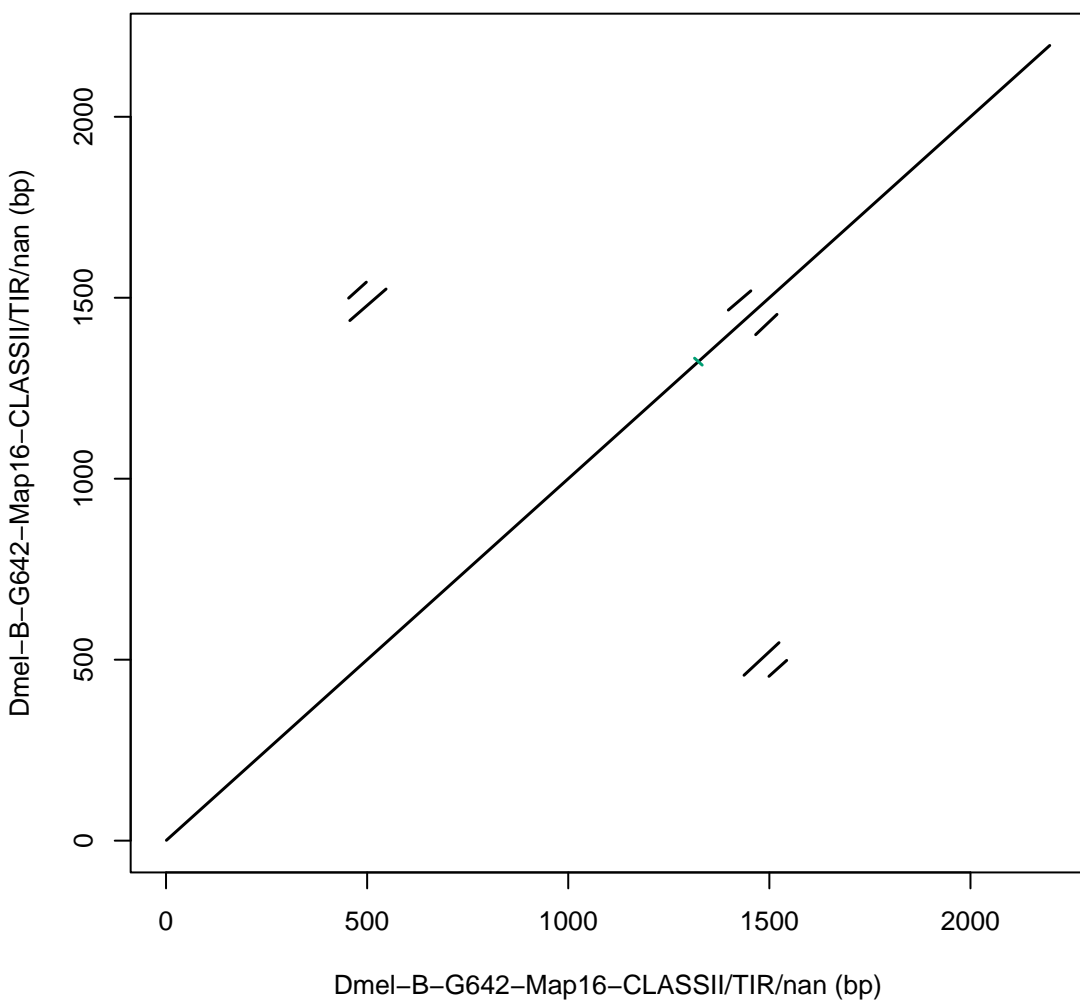**TE: Dmel−B−G642−Map16−CLASSII/TIR/nan**
consensus size: 2197bp; fragments: 107; full length: 1 (>=1977.3bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected