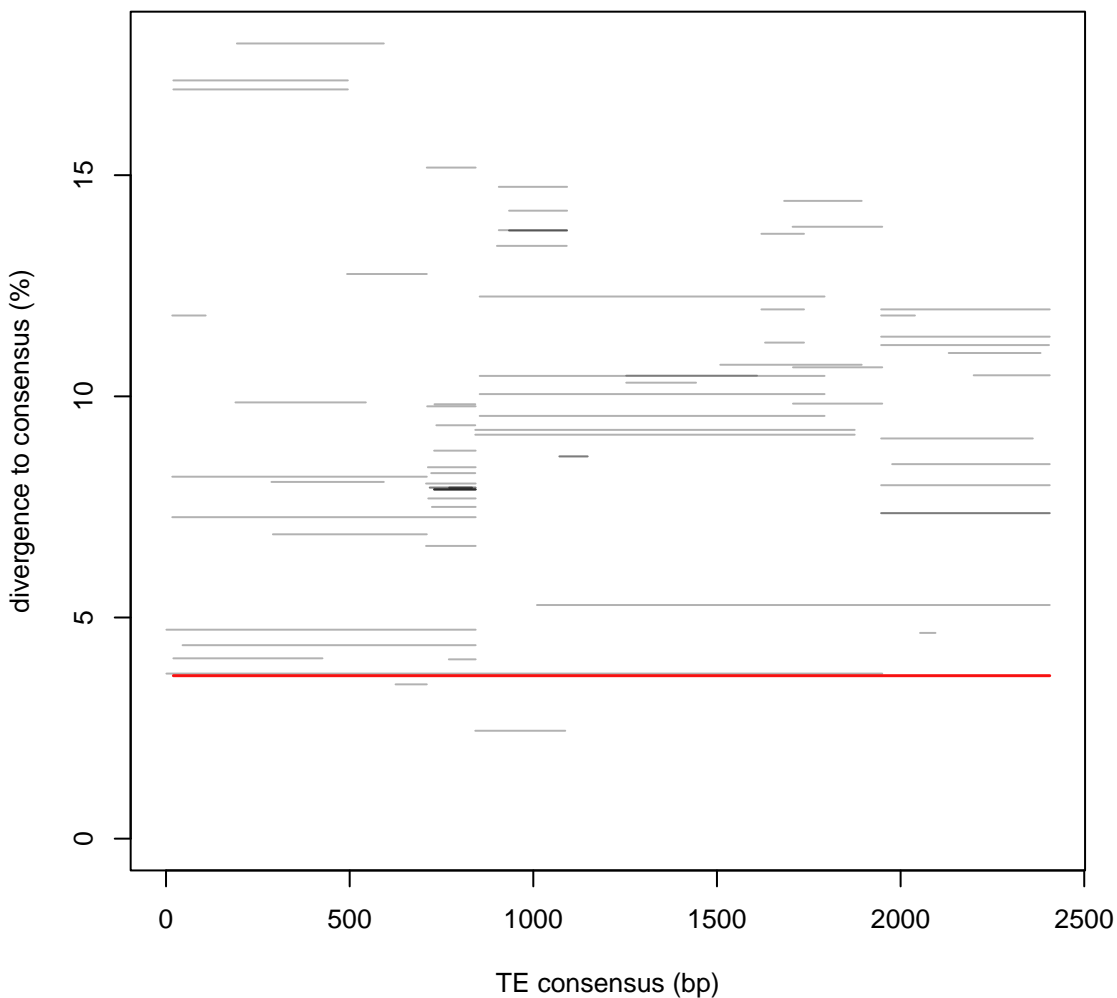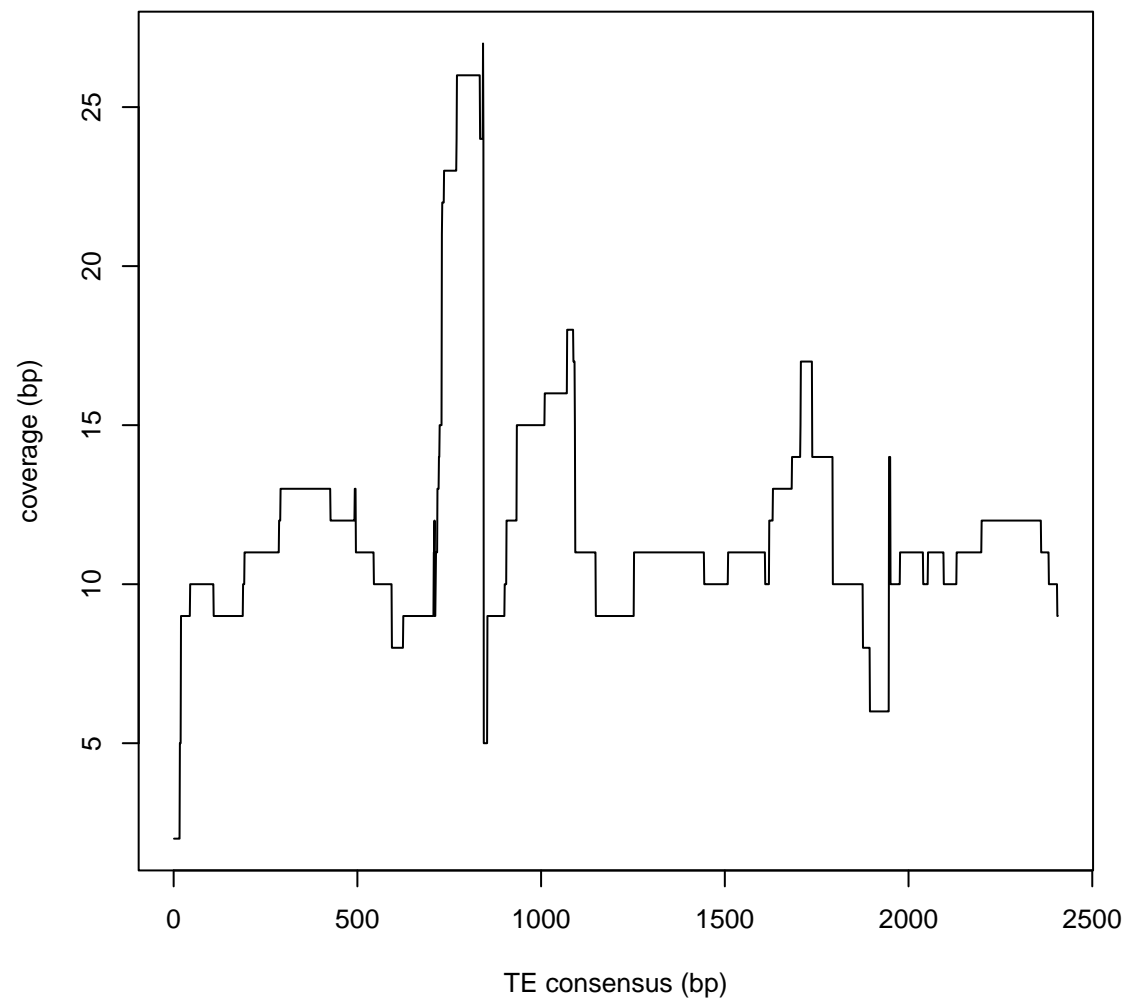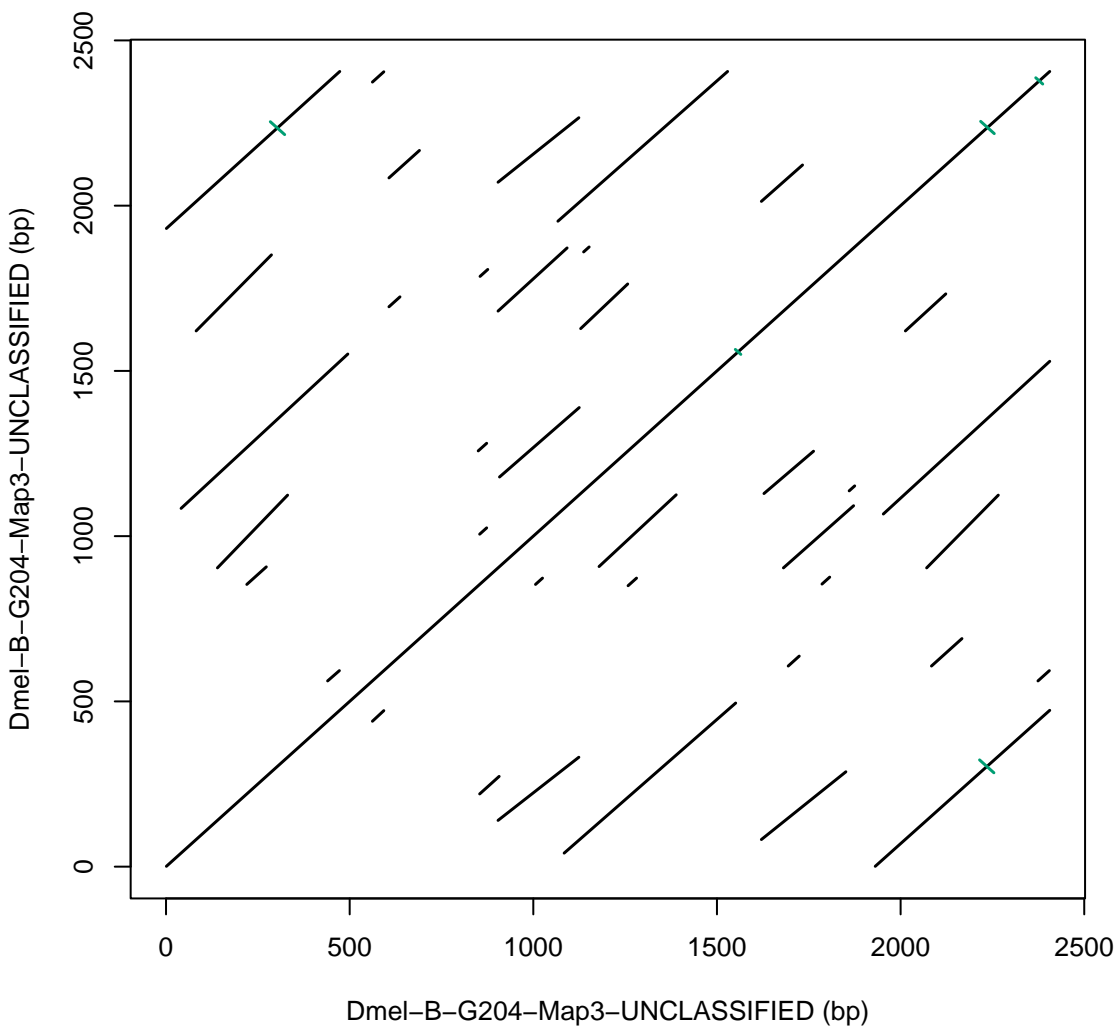**TE: Dmel−B−G204−Map3−UNCLASSIFIED**
consensus size: 2406bp; fragments: 75; full length: 1 (>=2165.4bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected