TE: Dmel-B-R884-Map3-TIR/nan
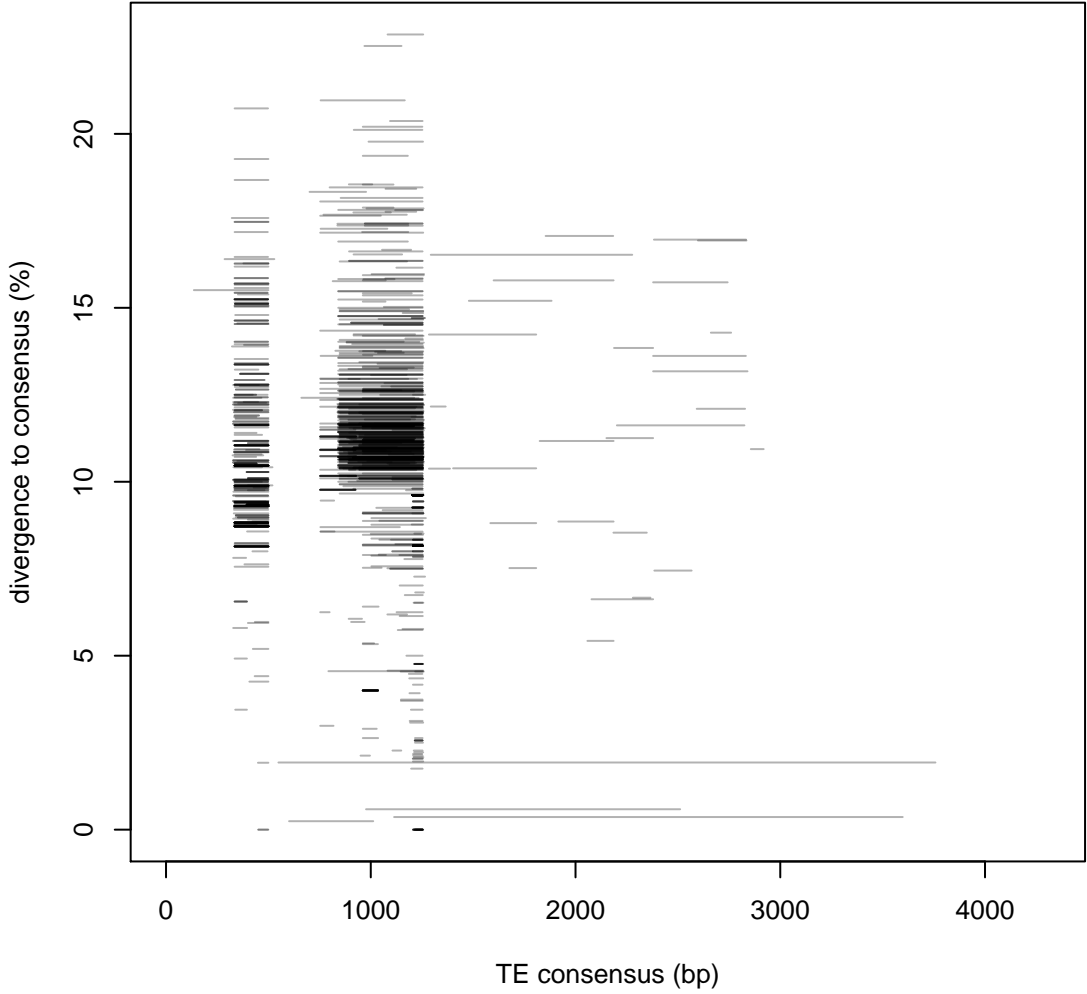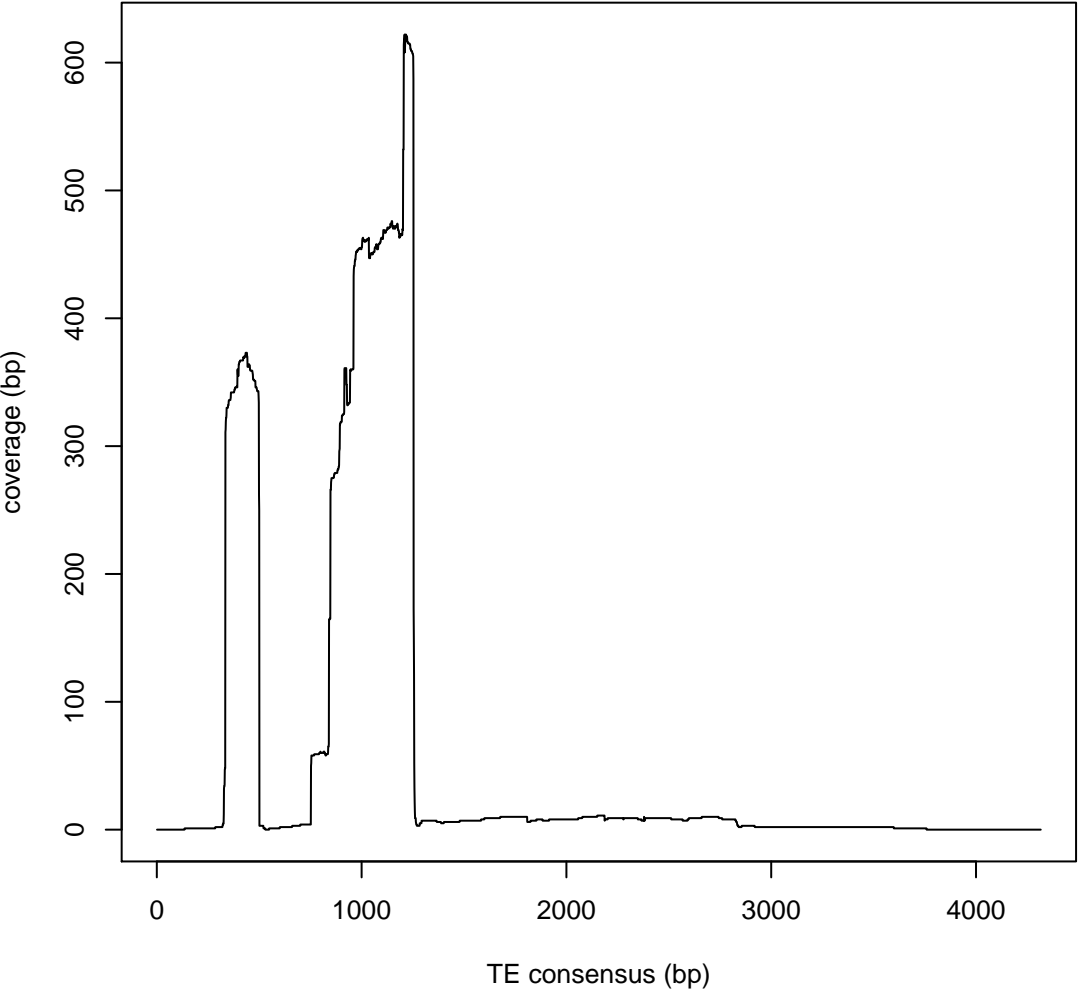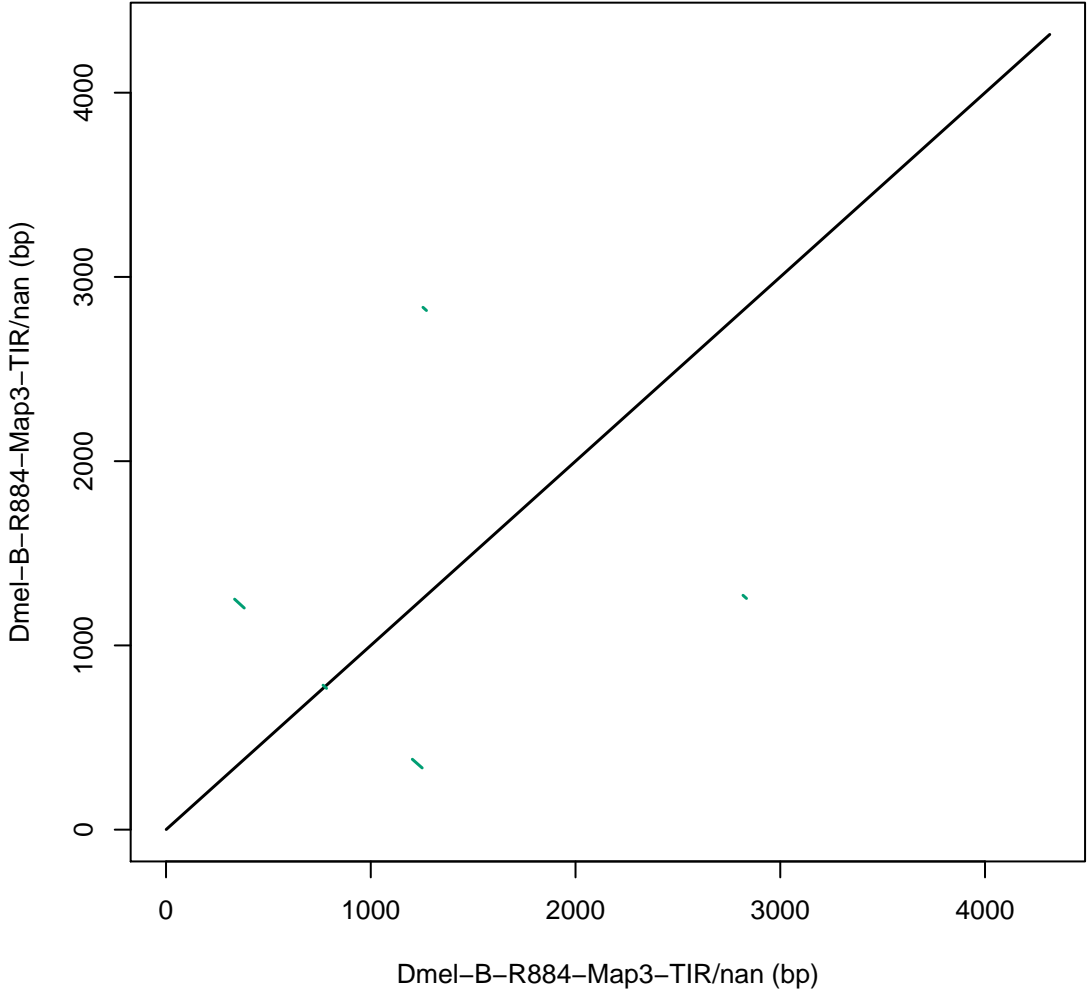consensus size: 4316bp; fragments: 1162; full length: 0 (>=3884.4bp)

divergence to consensus (%)

TE consensus (bp)

TE consensus genomic coverage

coverage (bp)

TE consensus (bp)

TE consensus self dotplot (blastn)

Dmel-B-R884-Map3-TIR/nan (bp)

Dmel-B-R884-Map3-TIR/nan (bp)

TE consensus structure and protein hits

no orf >400 bp detected

Dmel-B-R884-Map3-TIR/nan (bp)