TE: Dmel−B−P70.91−Map20−TIR/nan
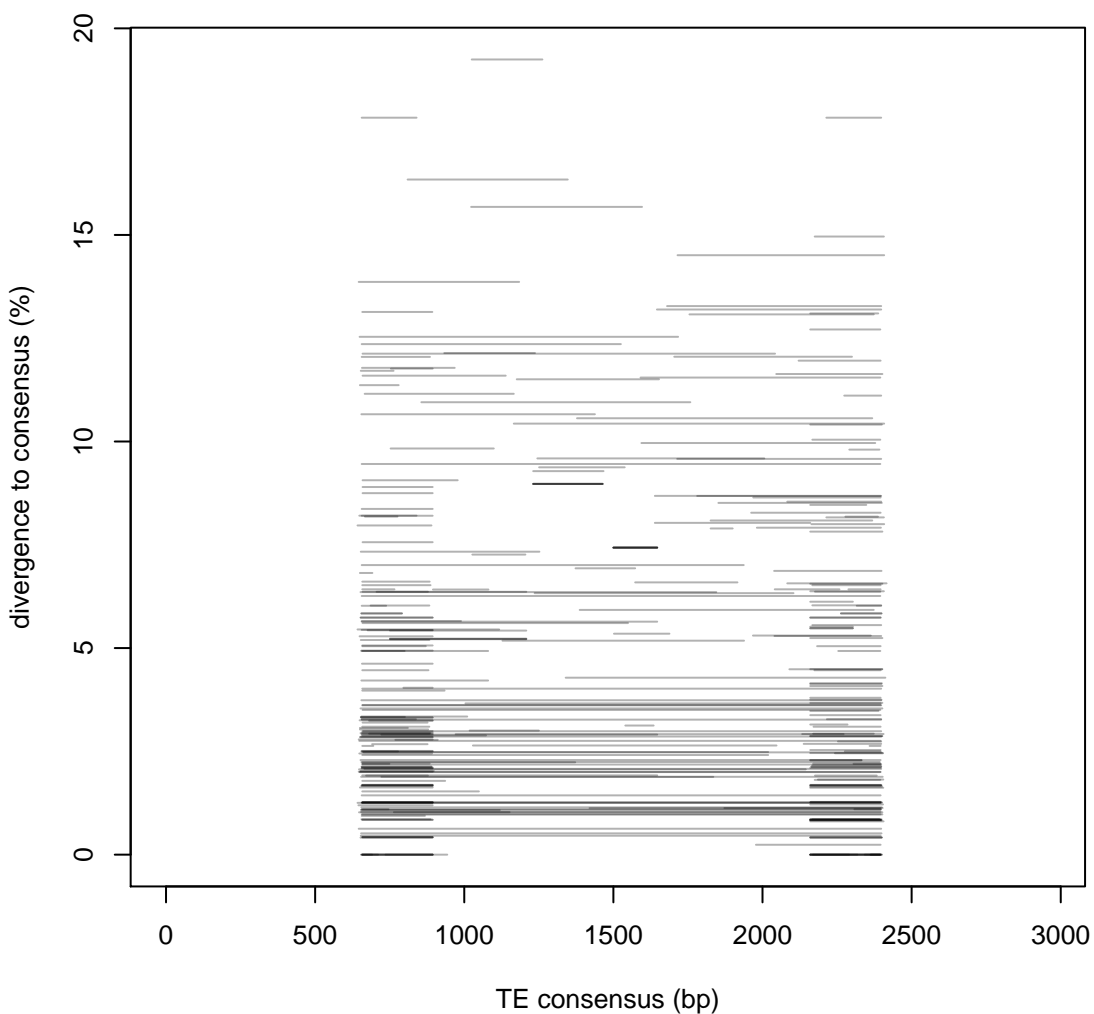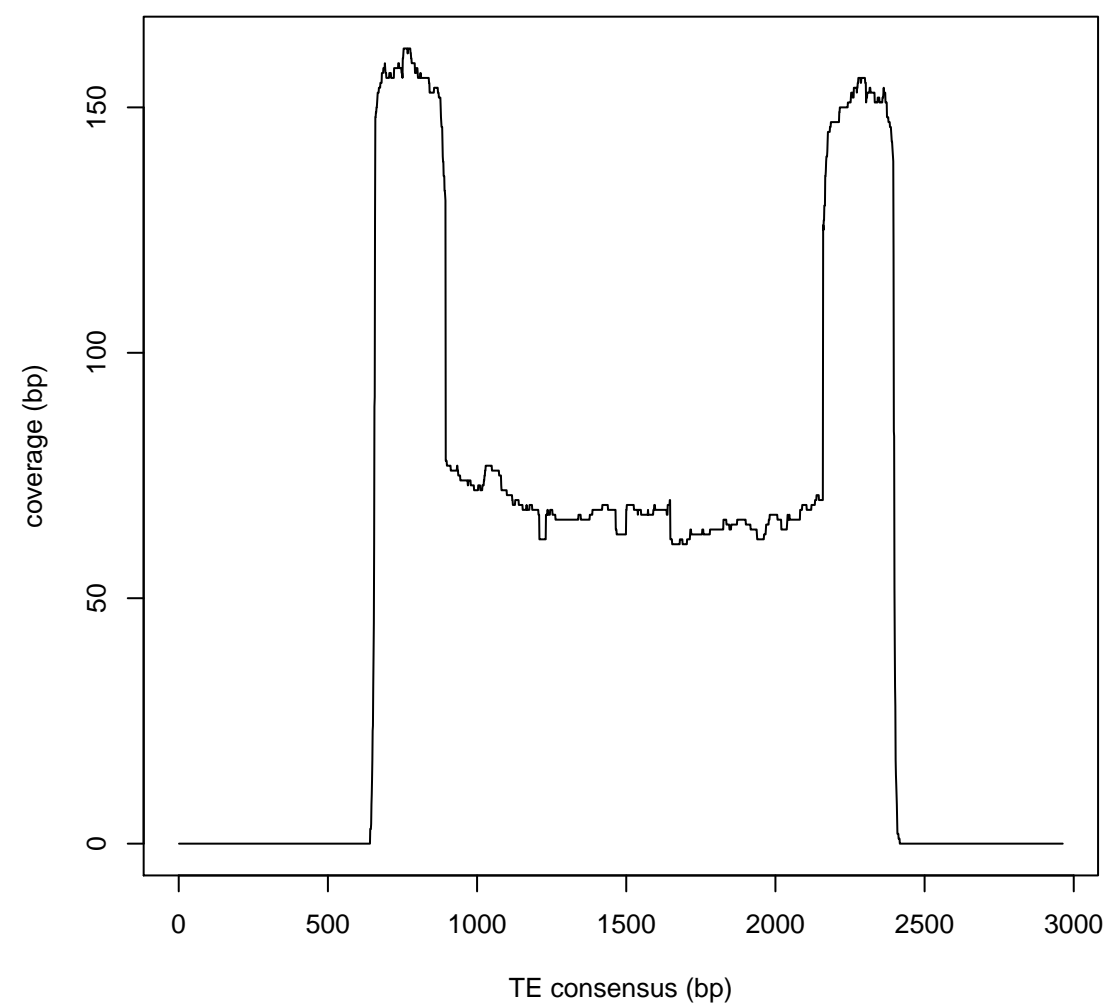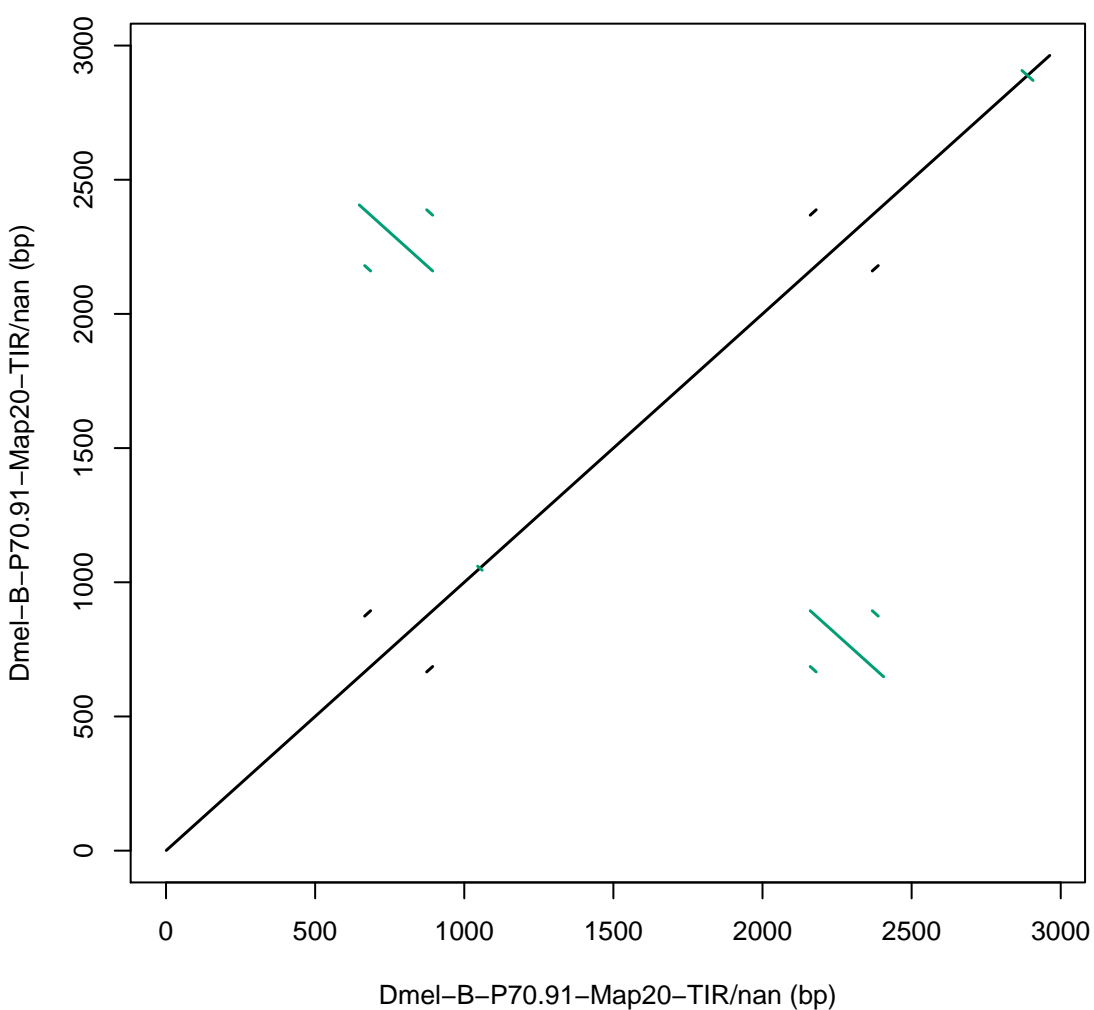consensus size: 2963bp; fragments: 335; full length: 0 (>=2666.7bp)

TE consensus genomic coverage

divergence to consensus (%)

TE consensus (bp)

coverage (bp)

TE consensus (bp)

TE consensus self dotplot (blastn)

Dmel−B−P70.91−Map20−TIR/nan (bp)

Dmel−B−P70.91−Map20−TIR/nan (bp)

TE consensus structure and protein hits

no orf >400 bp detected

Dmel−B−P70.91−Map20−TIR/nan (bp)