TE: Dmel−B−R1598−Map6−Unclassified/nan
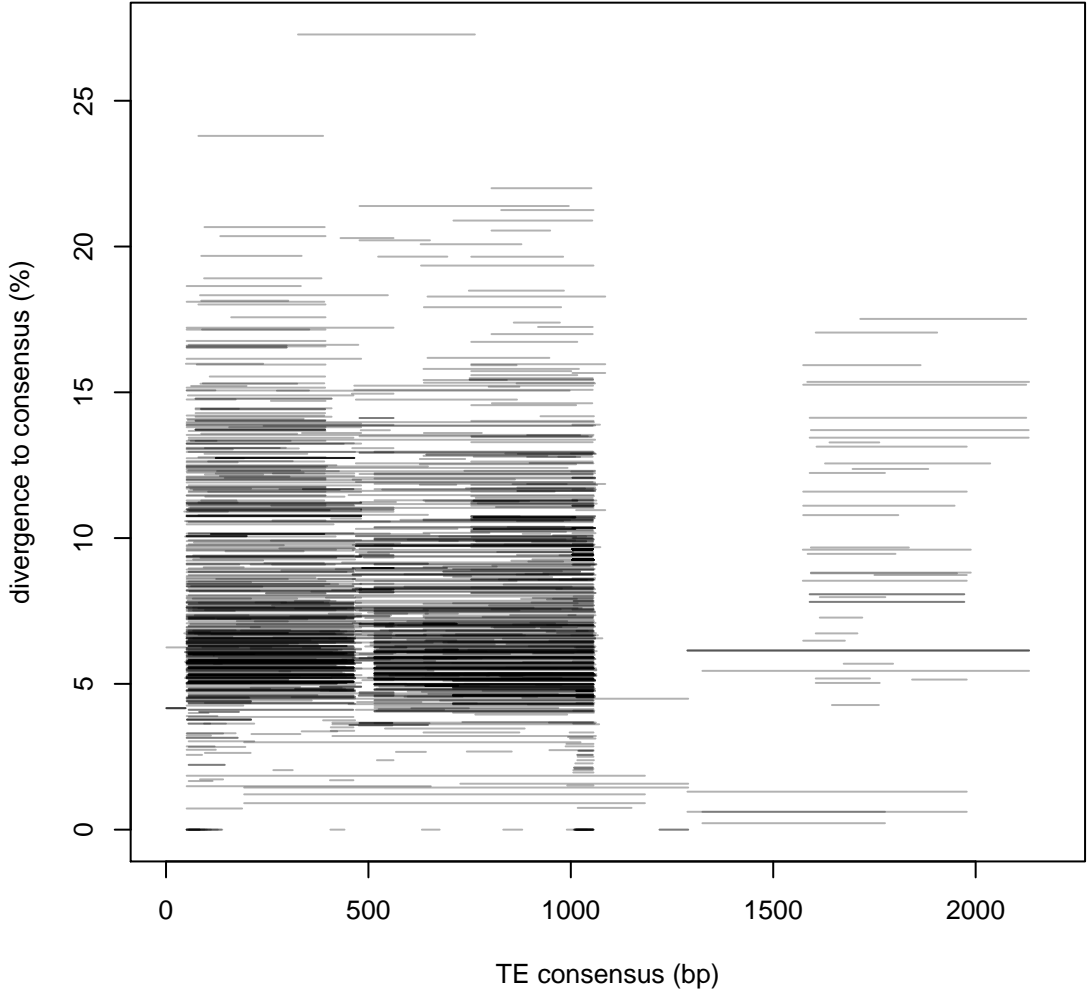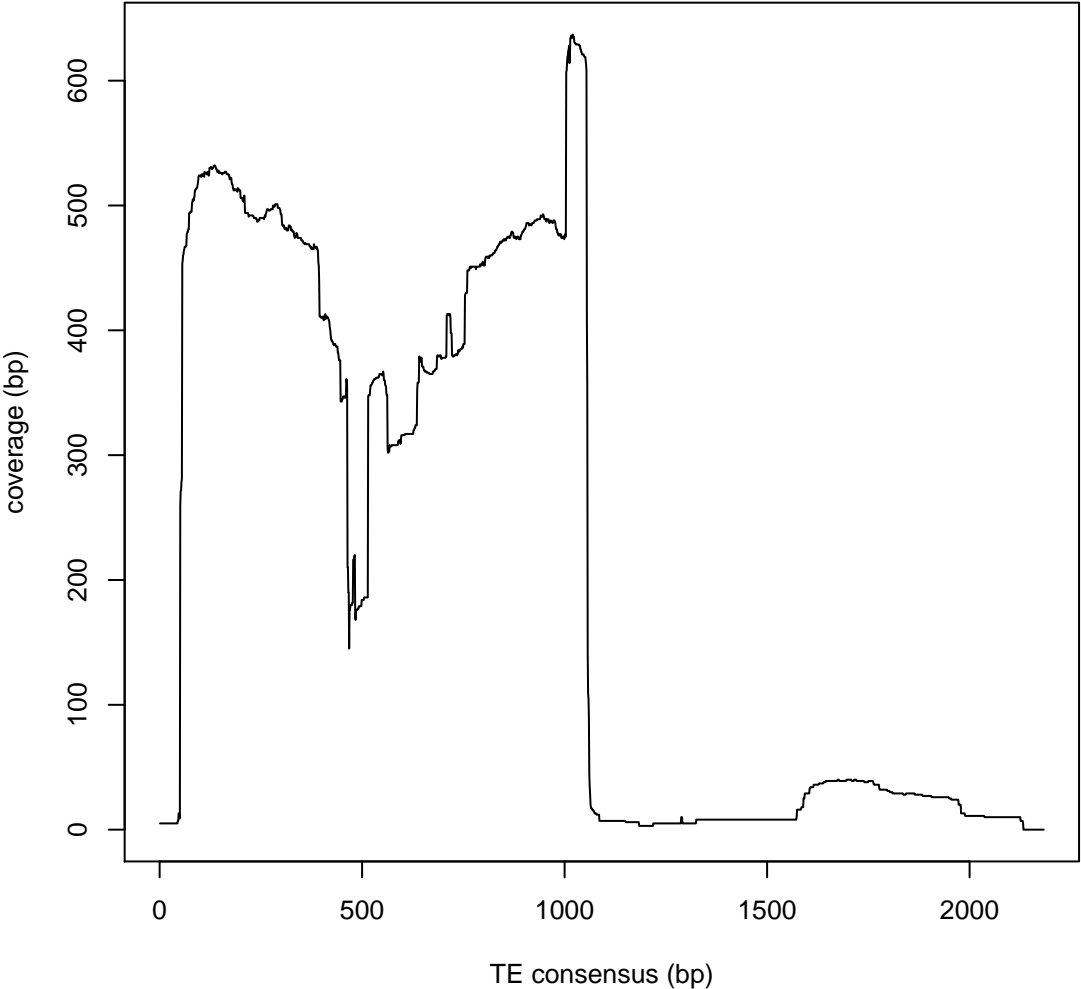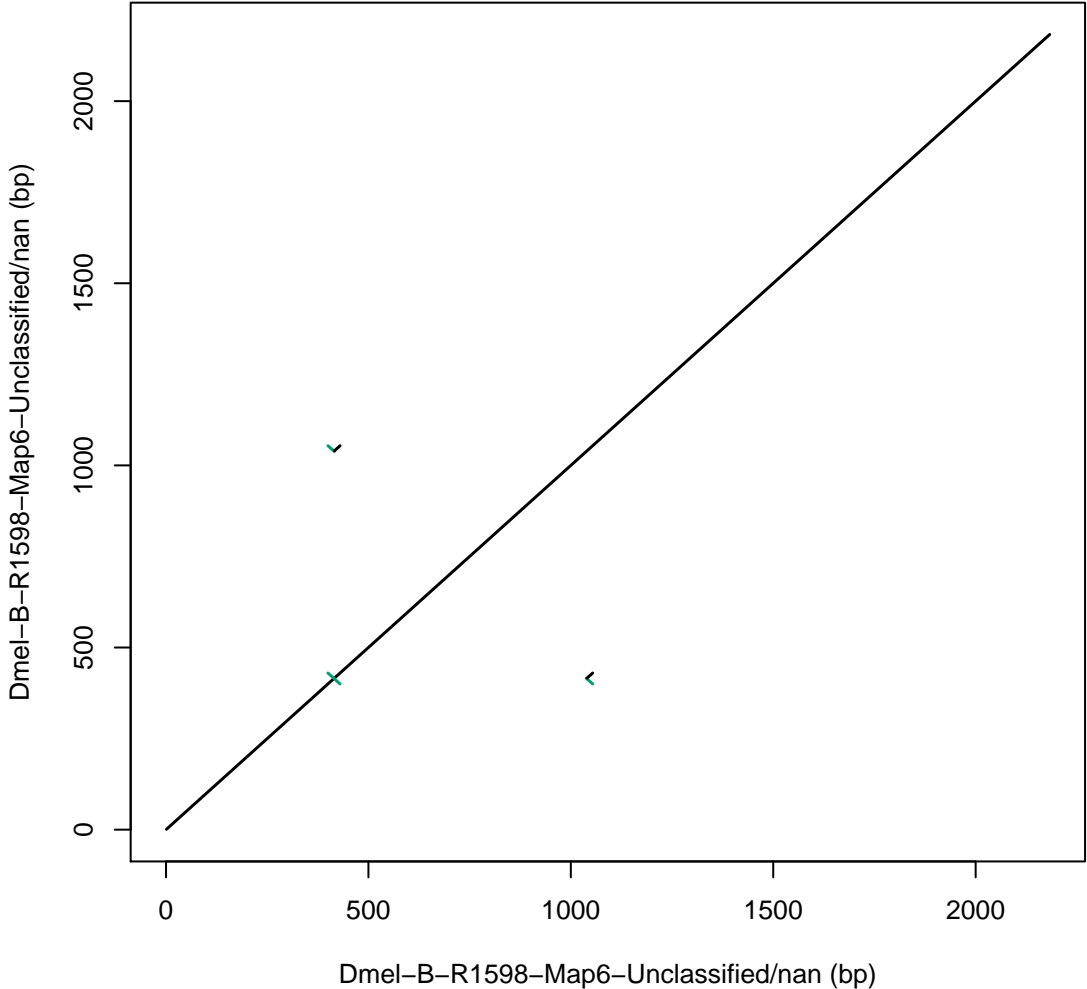**consensus size: 2183bp; fragments: 1493; full length: 0 (>=1964.7bp)**

TE consensus genomic coverage

divergence to consensus (%)

TE consensus (bp)

coverage (bp)

TE consensus (bp)

TE consensus self dotplot (blastn)

Dmel−B−R1598−Map6−Unclassified/nan (bp)

Dmel−B−R1598−Map6−Unclassified/nan (bp)

TE consensus structure and protein hits

no orf >400 bp detected

Dmel−B−R1598−Map6−Unclassified/nan (bp)