TE: Dmel−B−R895−Map6−LTR/Gypsy
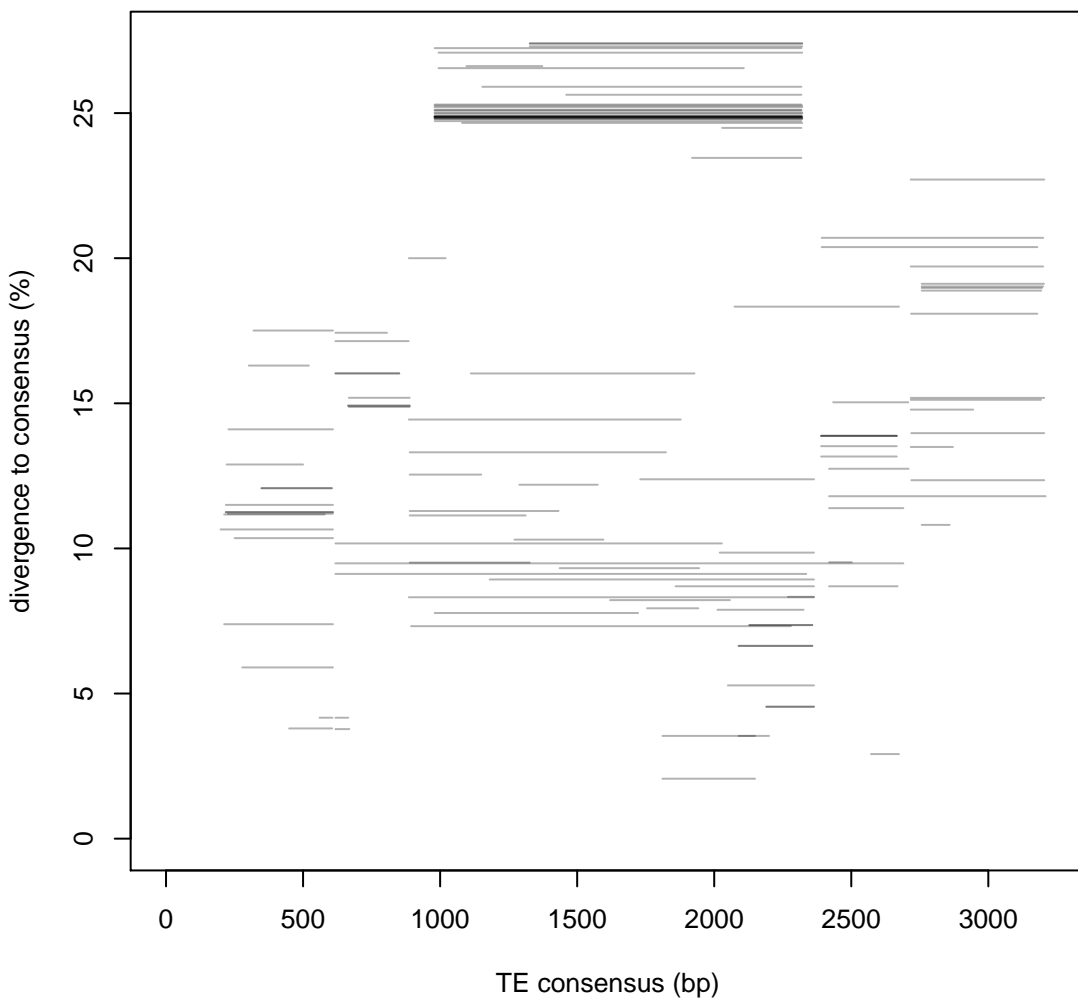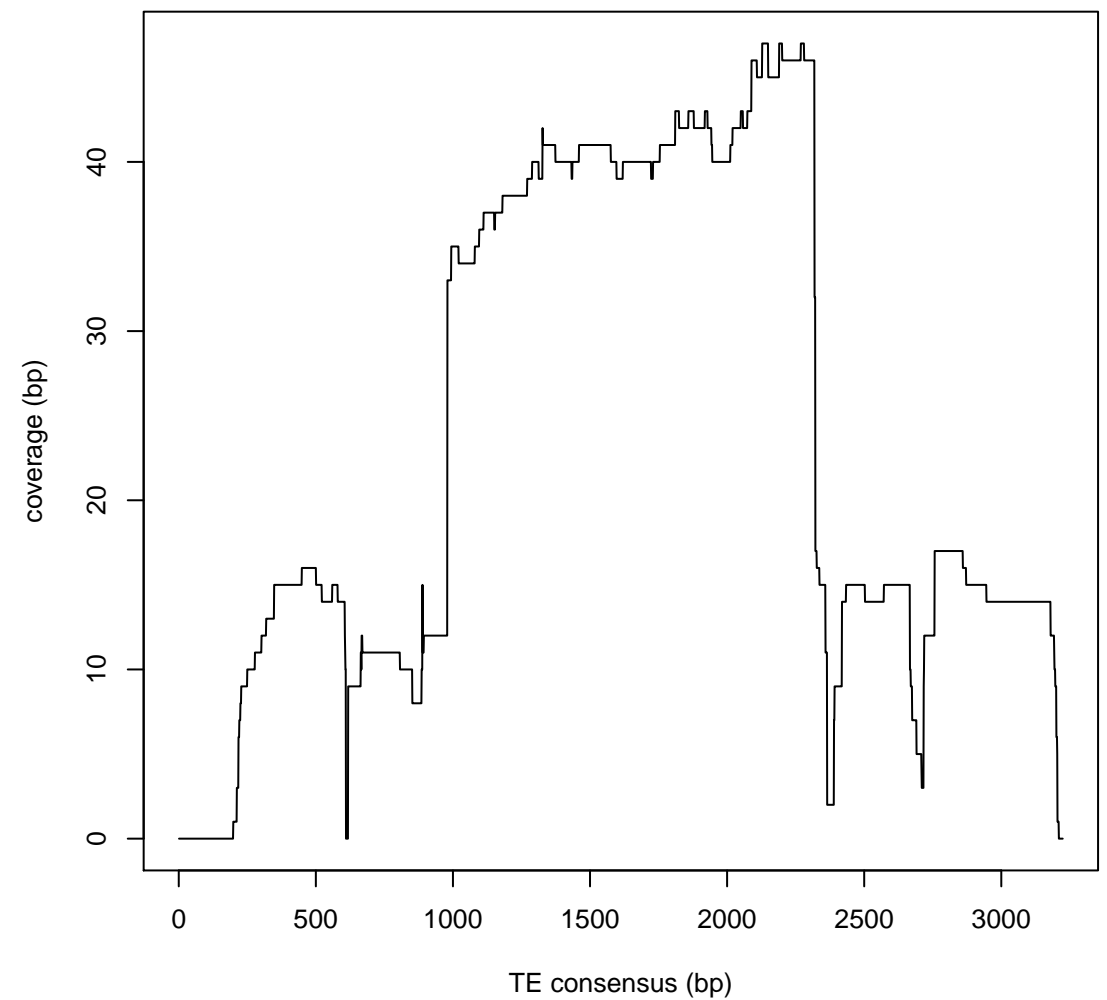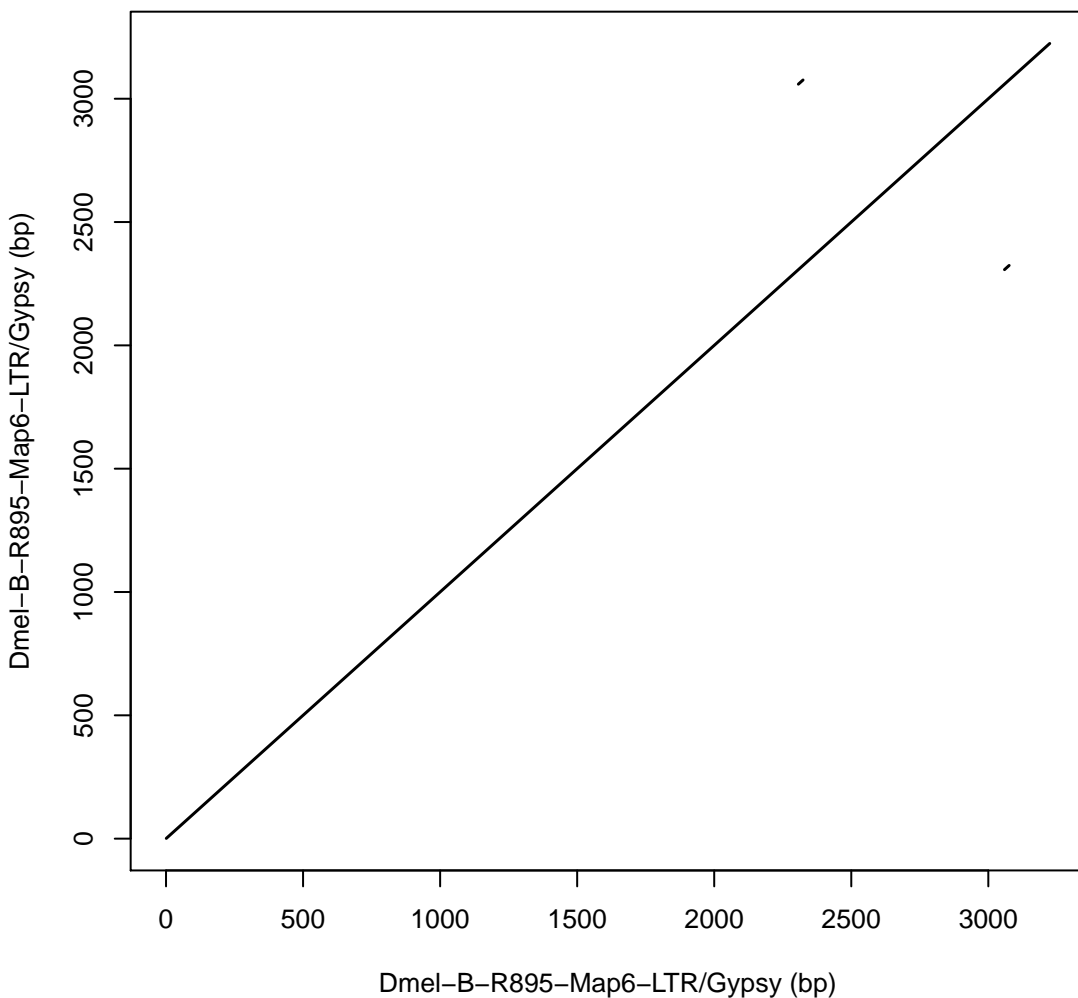**consensus size: 3224bp; fragments: 122; full length: 0 (>=2901.6bp)**

TE consensus genomic coverage

divergence to consensus (%)

TE consensus (bp)

coverage (bp)

TE consensus (bp)

TE consensus self dotplot (blastn)

Dmel−B−R895−Map6−LTR/Gypsy (bp)

Dmel−B−R895−Map6−LTR/Gypsy (bp)

TE consensus structure and protein hits

no orf >400 bp detected

Dmel−B−R895−Map6−LTR/Gypsy (bp)