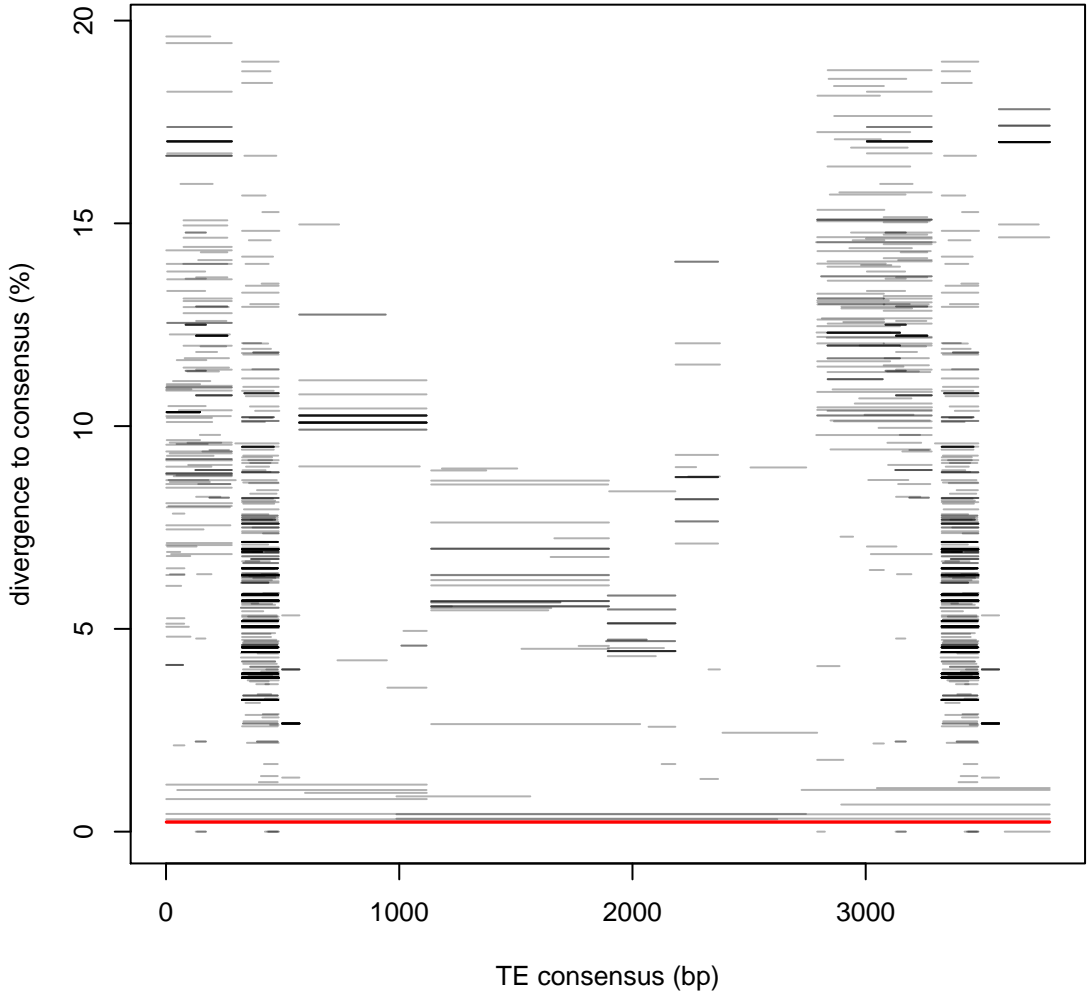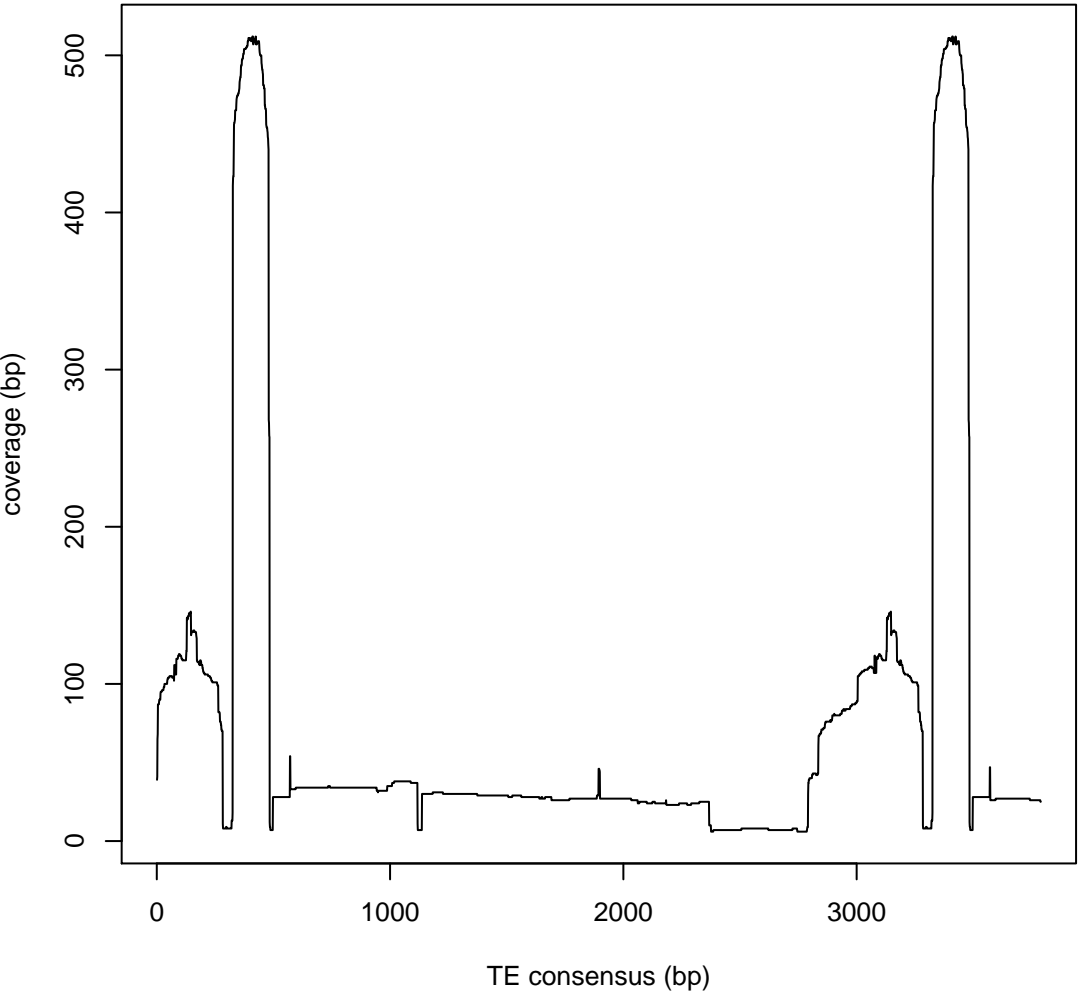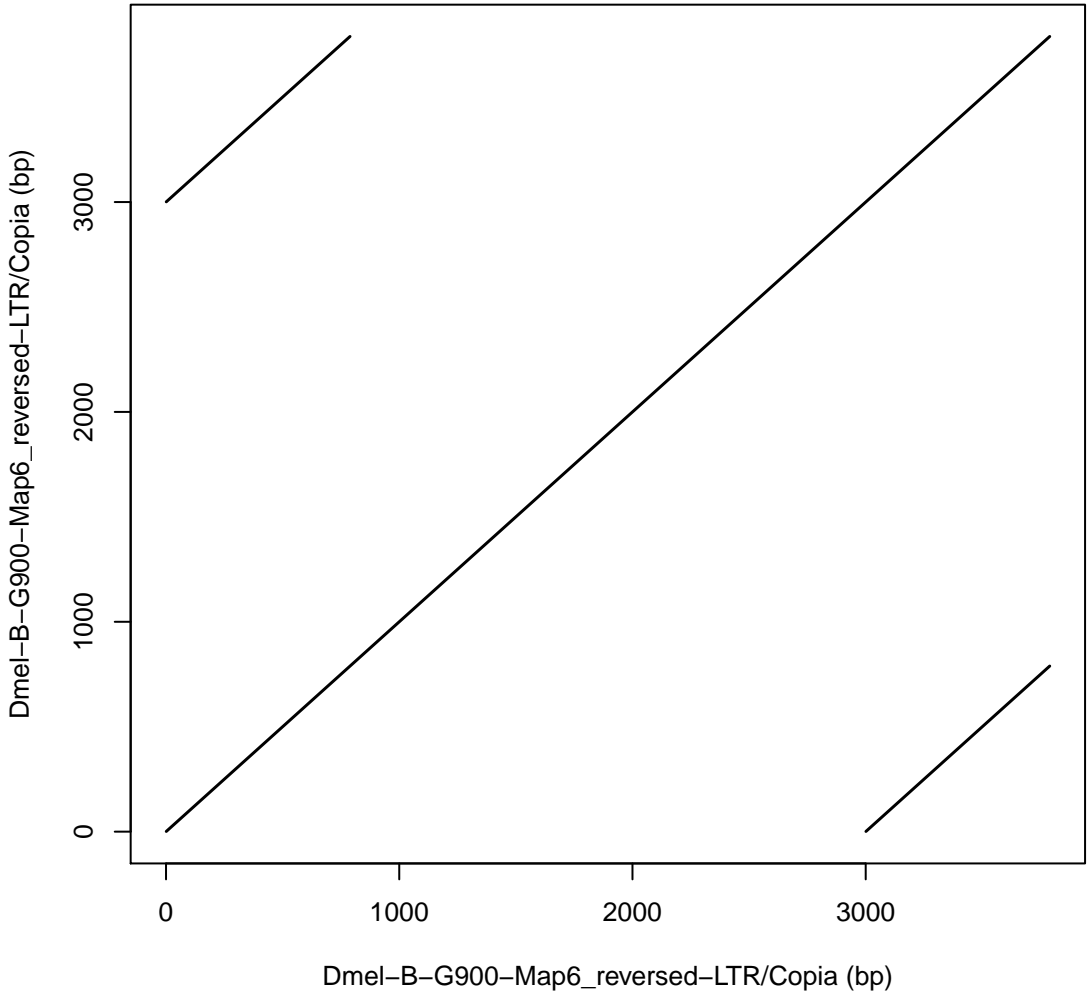**TE: DmeI−B−G900−Map6_reversed−LTR/Copia**
consensus size: 3789bp; fragments: 1559; full length: 2 (>=3410.1bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected