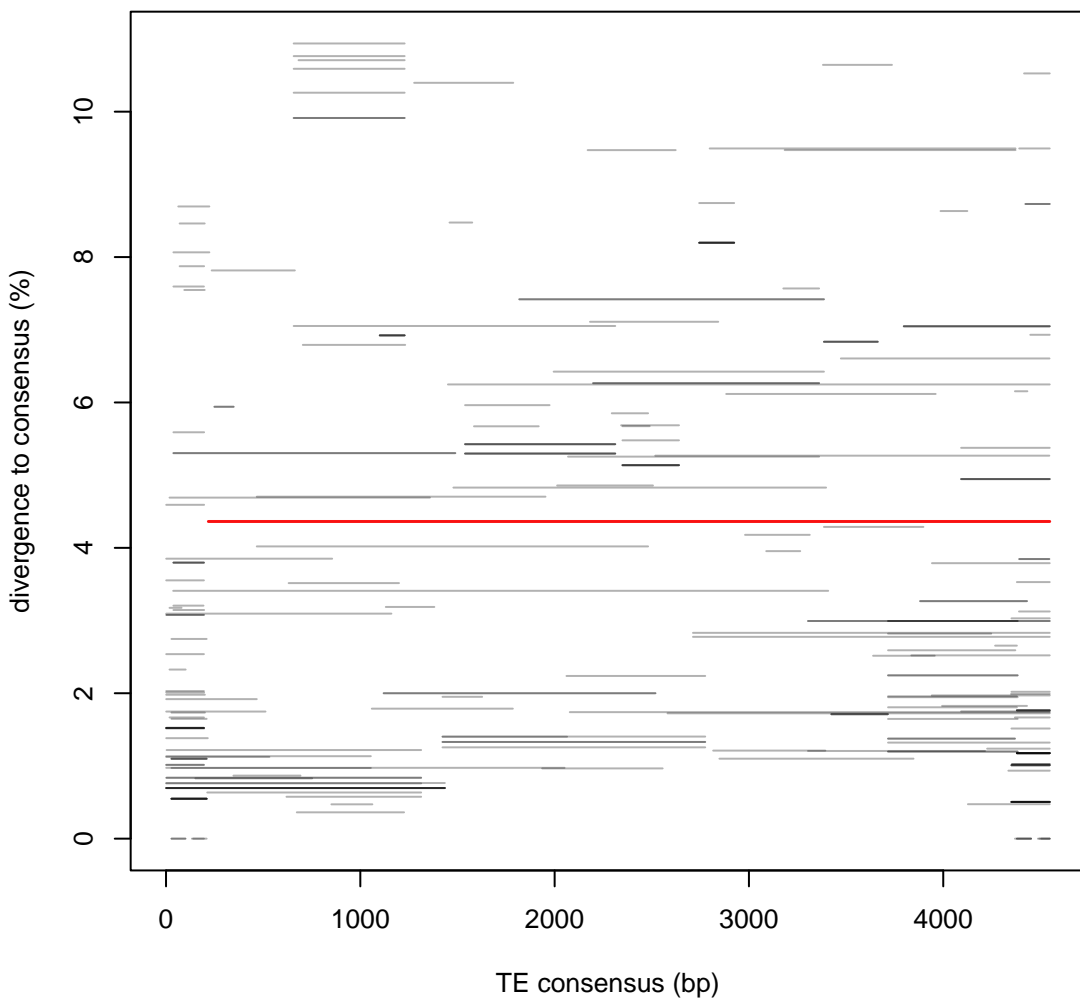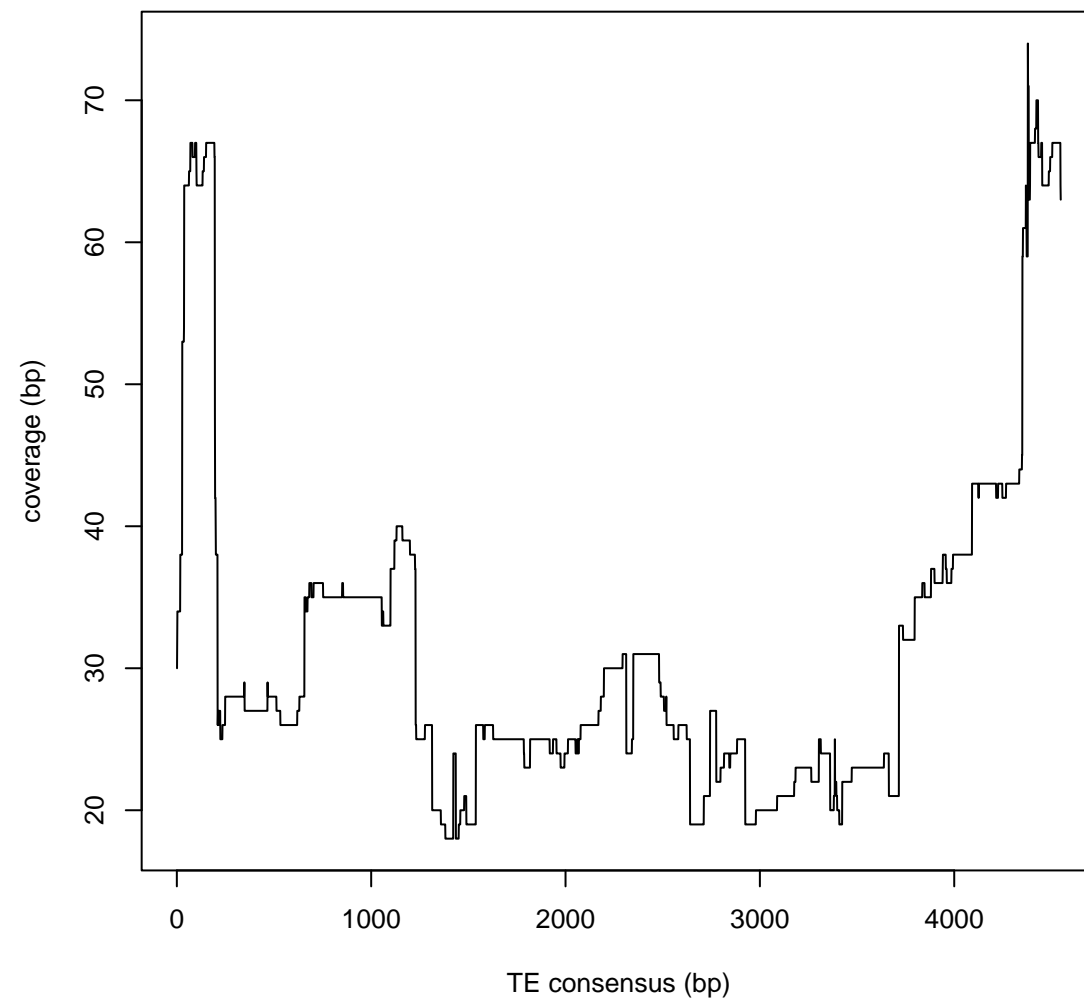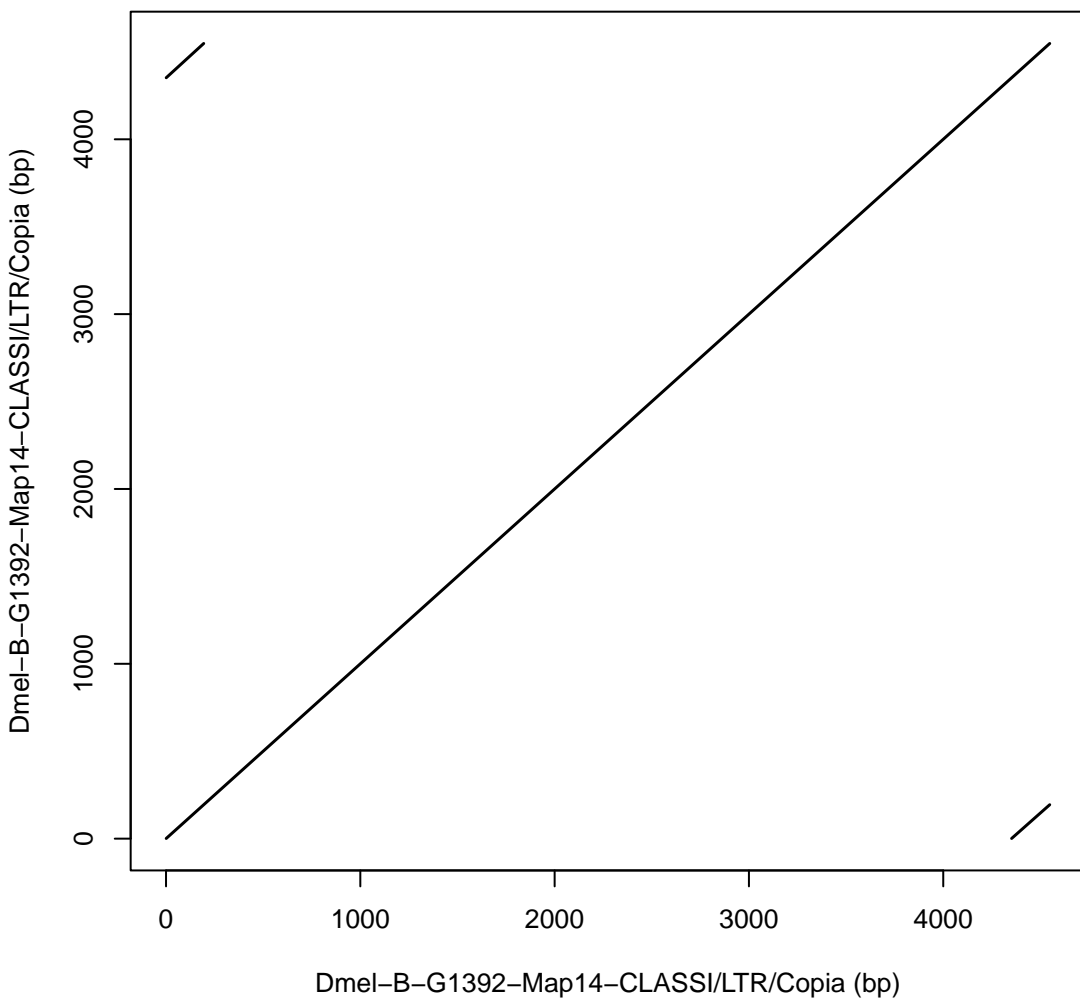**TE: DmeI−B−G1392−Map14−CLASSI/LTR/Copia**
consensus size: 4548bp; fragments: 249; full length: 1 (>=4093.2bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

Copia2 LTR/Copia

Copia2 LTR/Copia