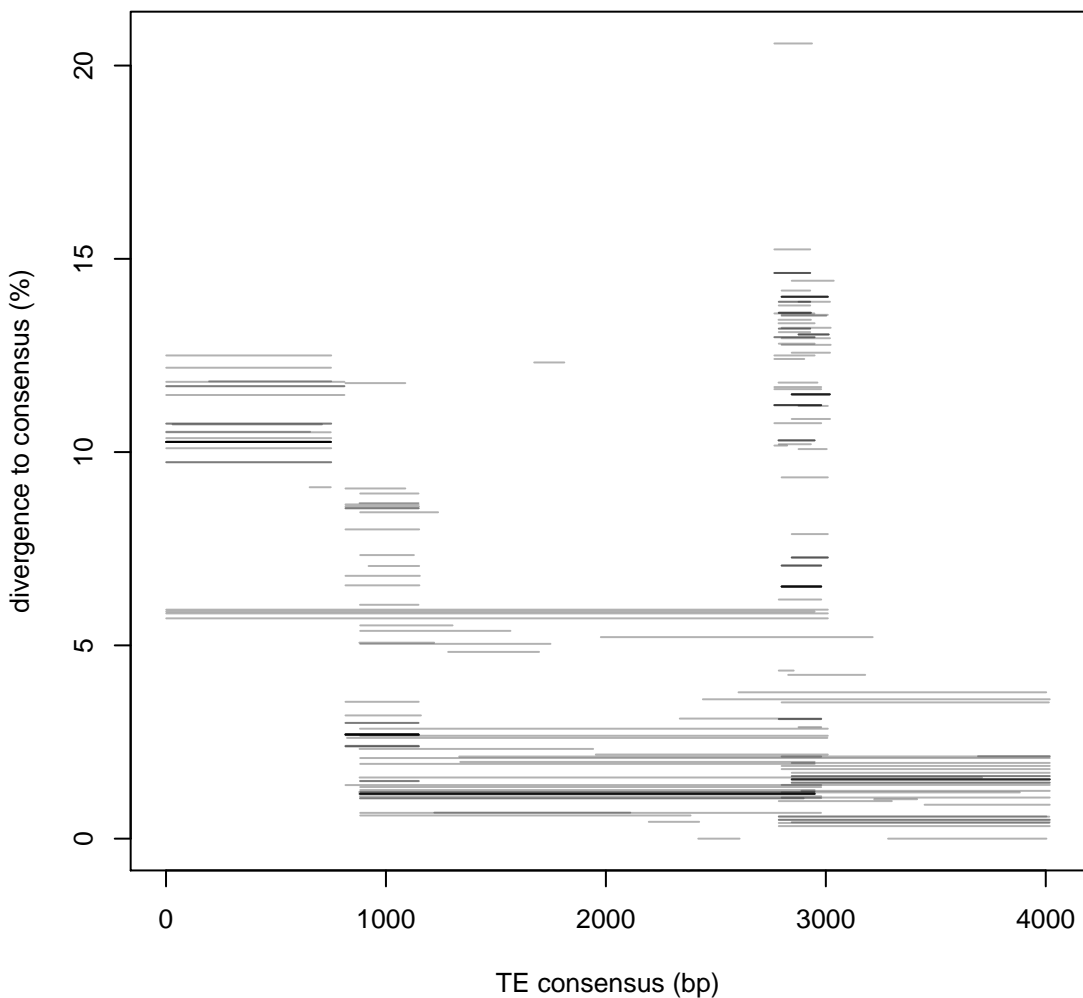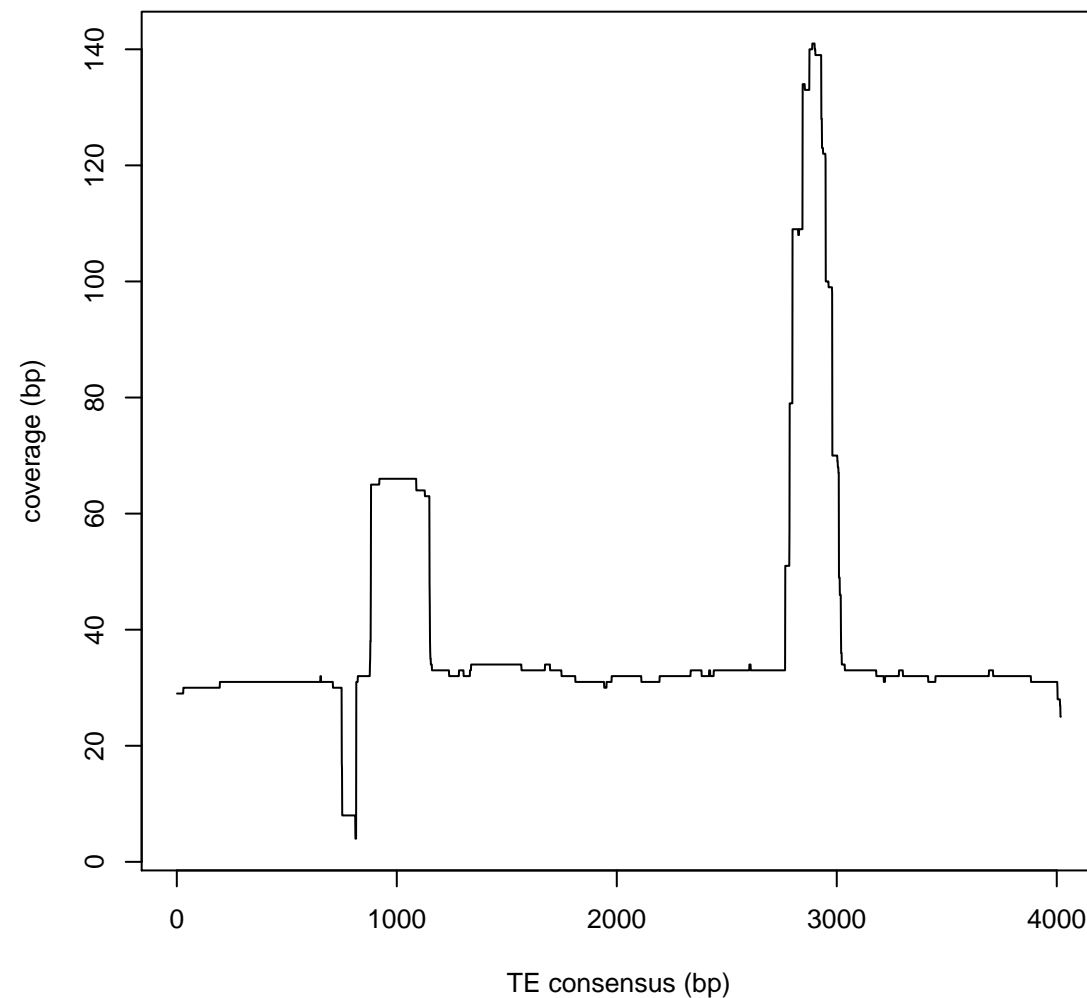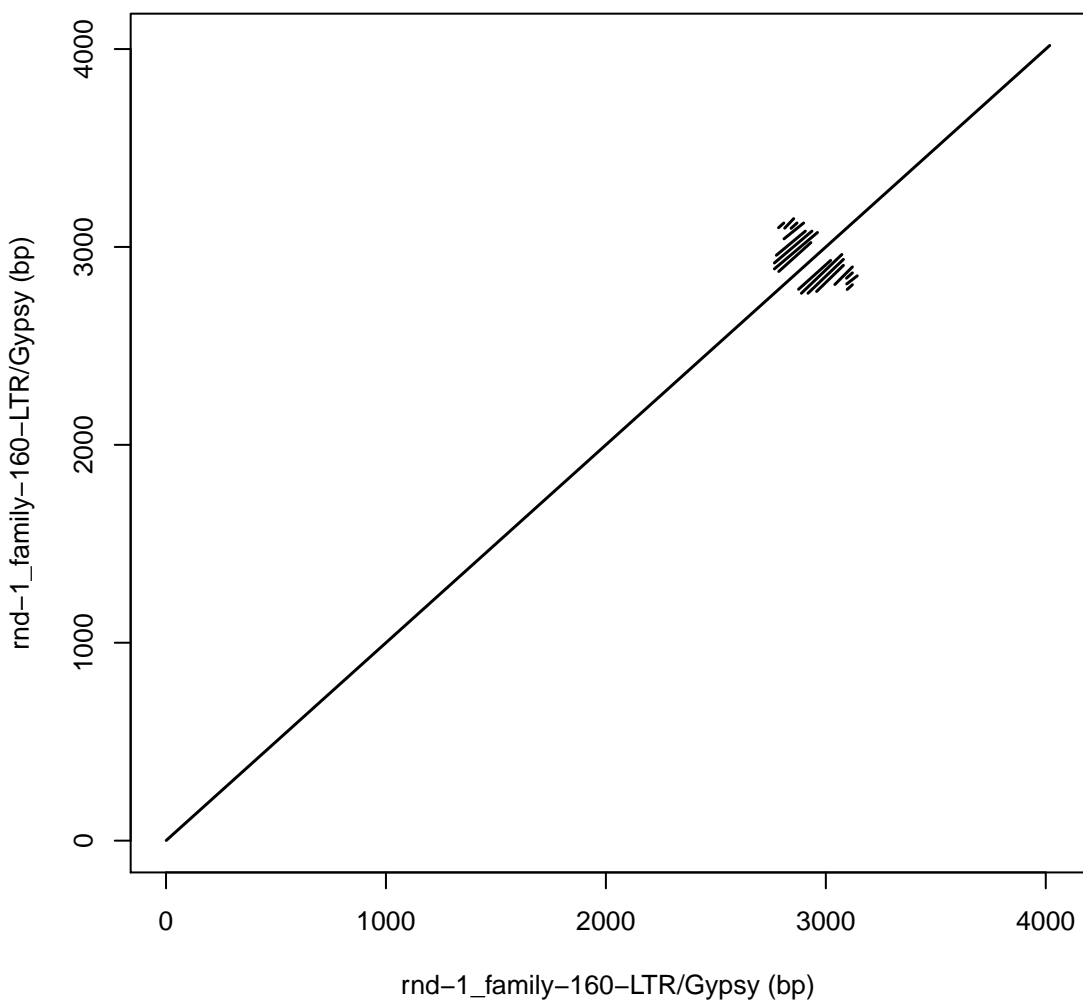TE: rnd−1_family−160−LTR/Gypsy
consensus size: 4018bp; fragments: 220; full length: 0 (>=3616.2bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

Copia−5_DAn LTR/Copia
BEL−5_DYa_pol LTI
MDG3 LTR/Gypsy