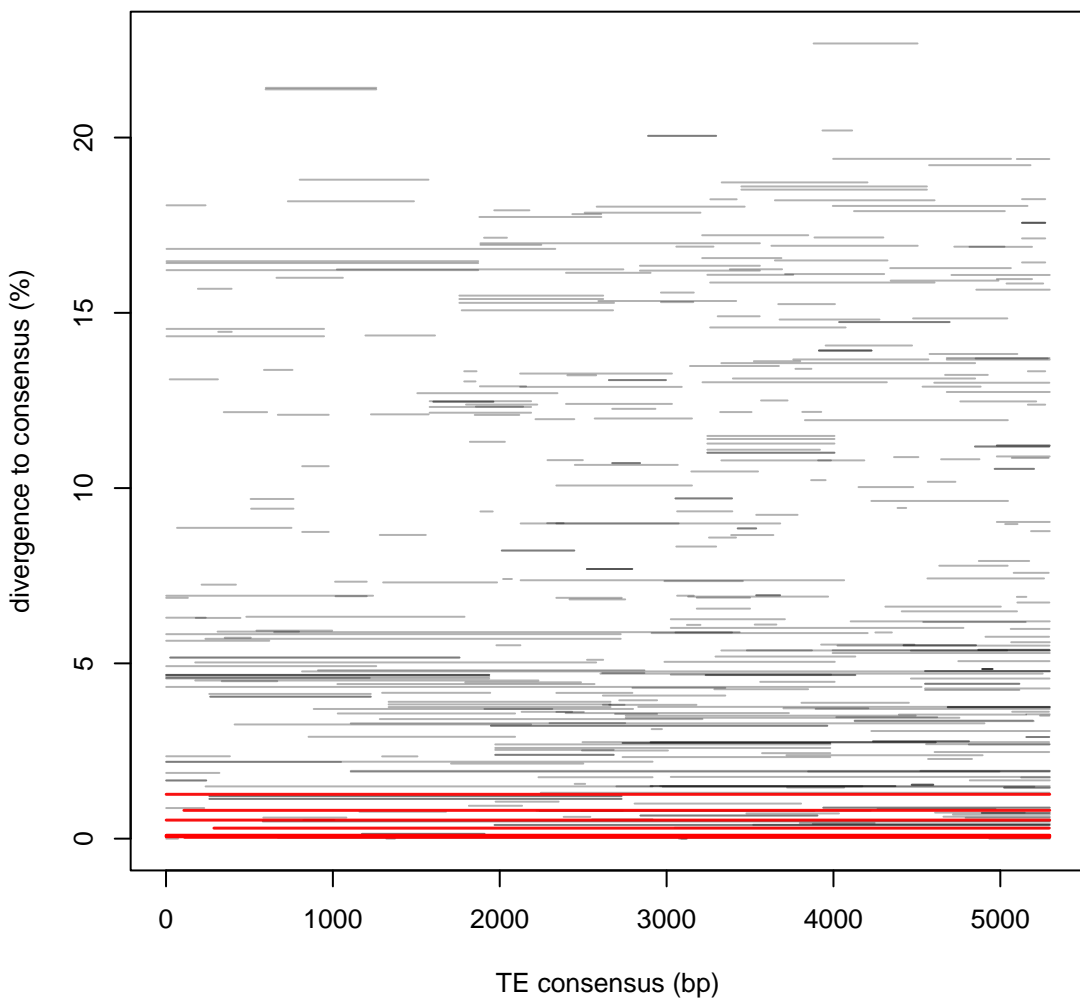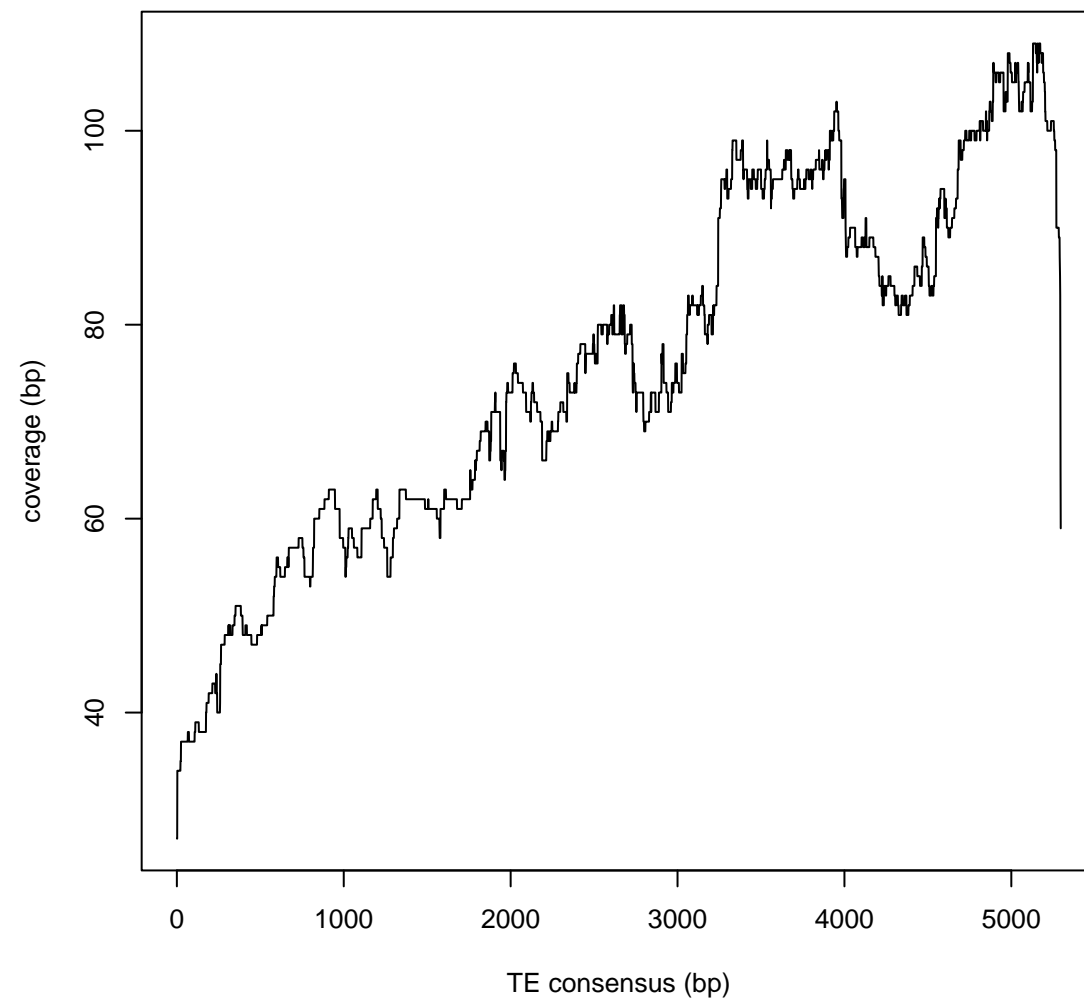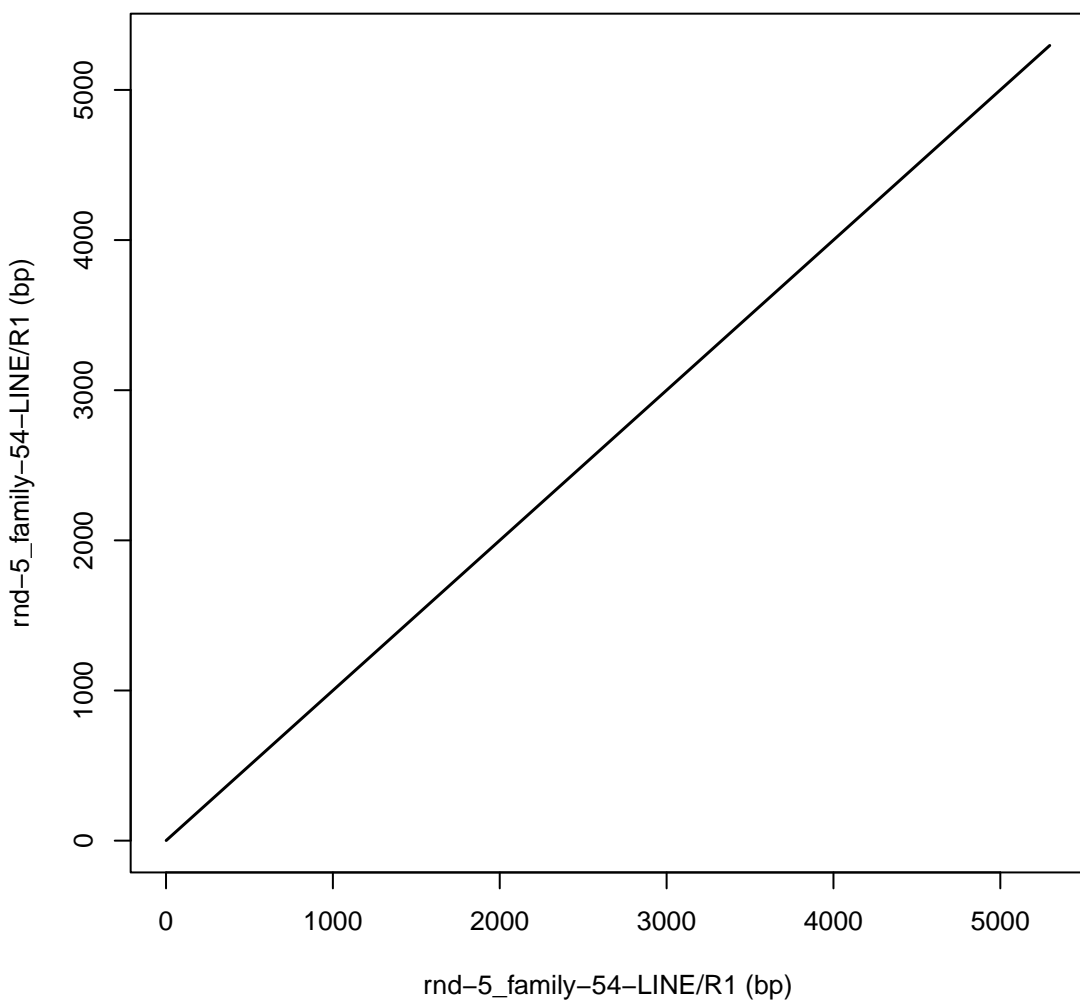**TE: rnd−5_family−54−LINE/R1**
consensus size: 5296bp; fragments: 444; full length: 9 (>=4766.4bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

divergence to consensus (%)

TE consensus (bp)

coverage (bp)

TE consensus (bp)

rnd−5_family−54−LINE/R1 (bp)

rnd−5_family−54−LINE/R1 (bp)

opia−14_Mad LTR/Copia

Copia−44_BG LTR/Copia

DMRT1B_gag LINE/R1

L1−13_CPa_pol LINE/L1

ATCopia77 LTR/Copia

Polinton−2_NV_py MAV/Maverick

DMRT1B_pol LINE/R1