TE: rnd−1_family−461−LINE/R1−LOA
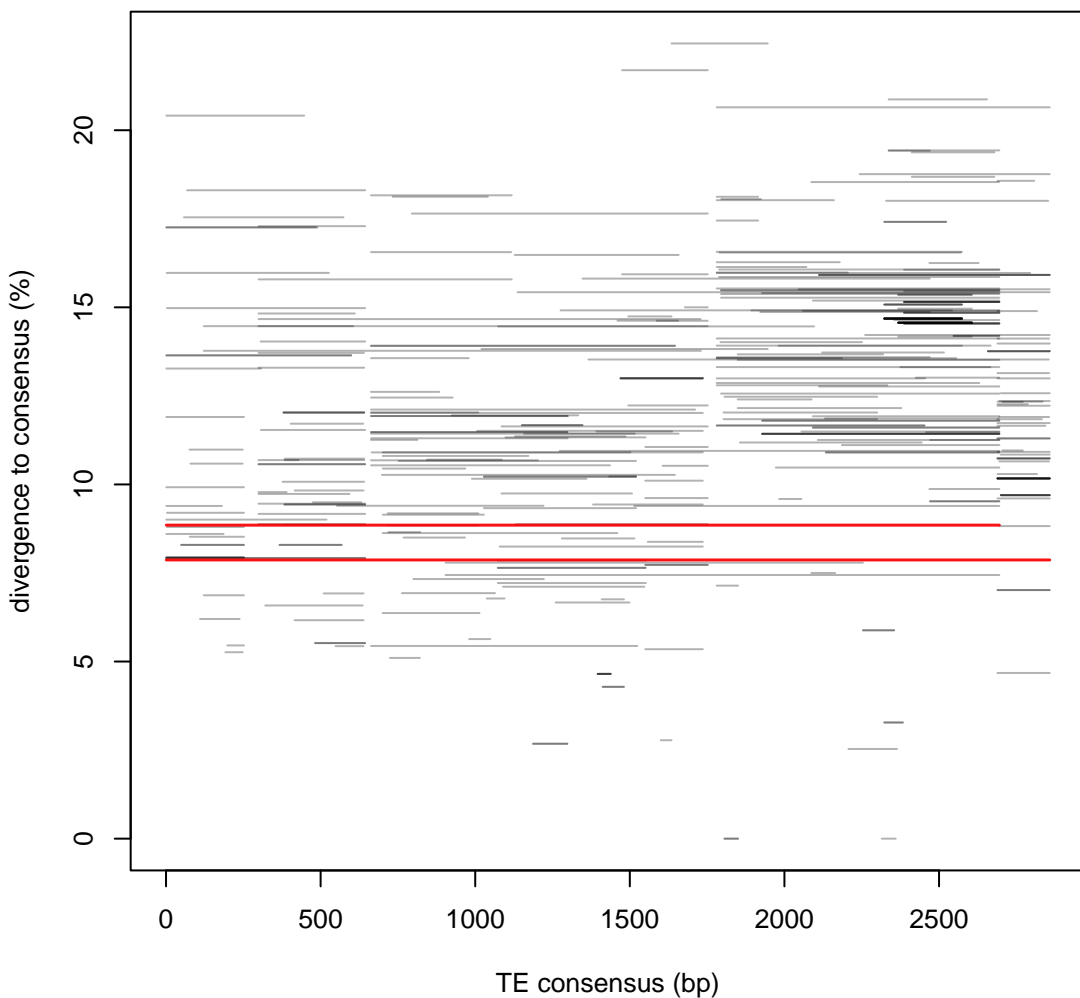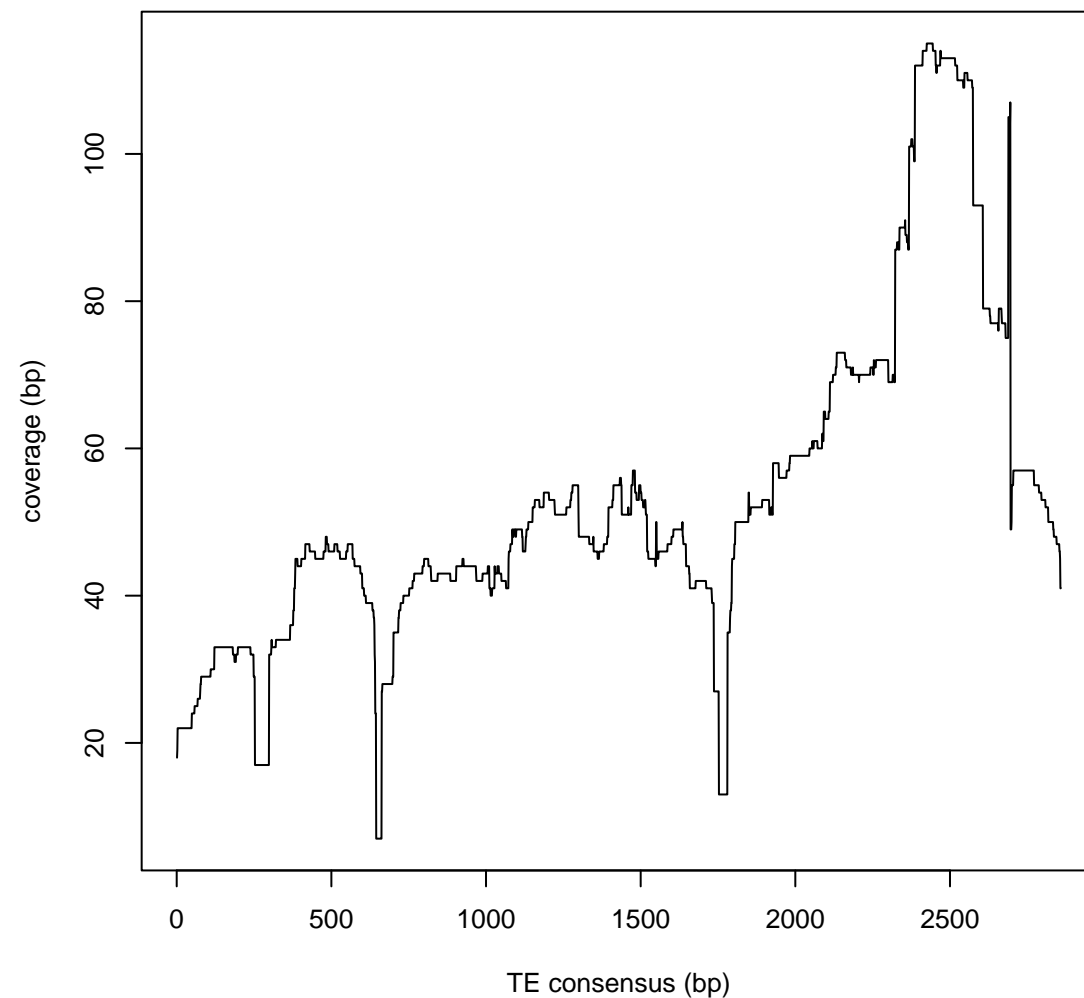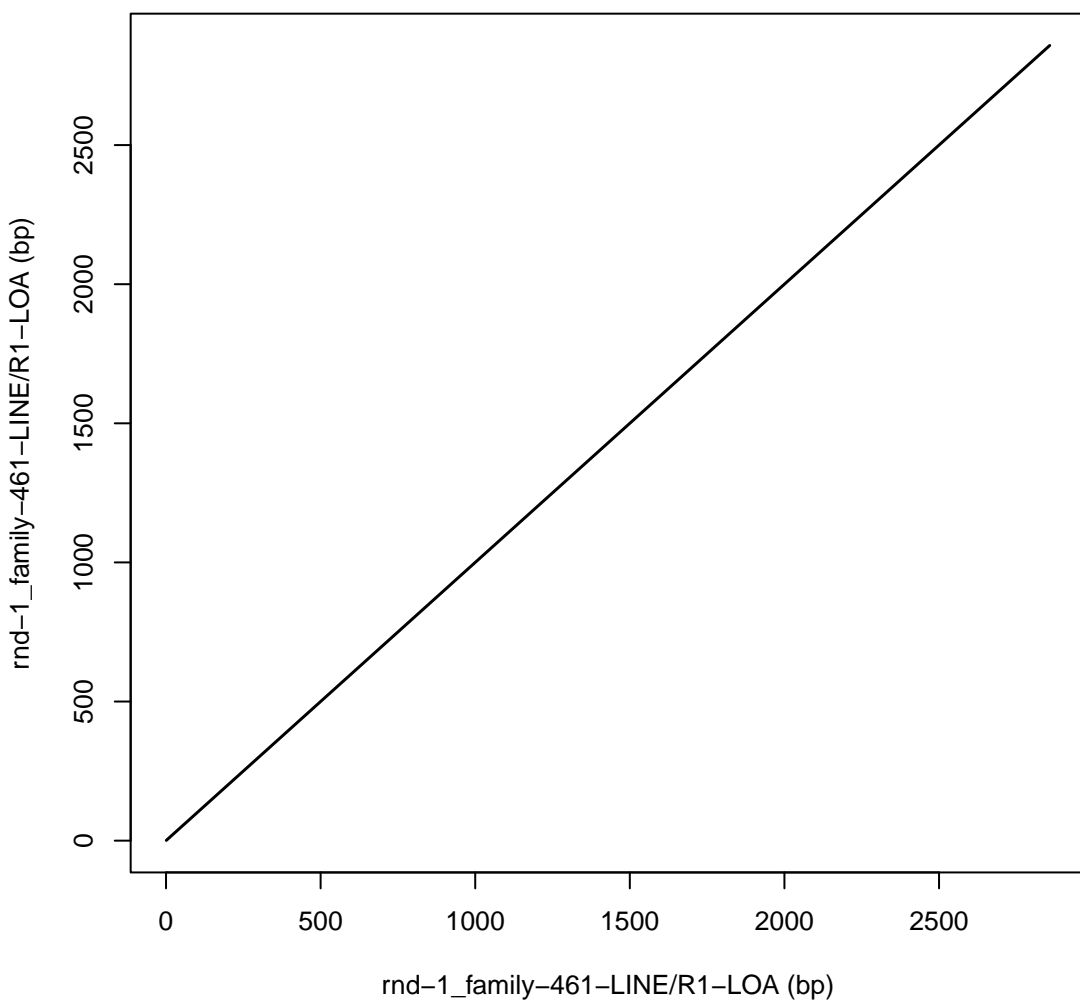**consensus size: 2858bp; fragments: 366; full length: 2 (>=2572.2bp)**

TE consensus (bp)

divergence to consensus (%)

TE consensus genomic coverage

coverage (bp)

TE consensus (bp)

TE consensus self dotplot (blastn)

rnd−1_family−461−LINE/R1−LOA (bp)

rnd−1_family−461−LINE/R1−LOA (bp)

TE consensus structure and protein hits

LOA_Ele6_pol LINE/R1−LOA

Crack−1_AFC_pol LINE/L2

rnd−1_family−461−LINE/R1−LOA (bp)