TE: rnd−5_family−1392−LTR/GYPSY
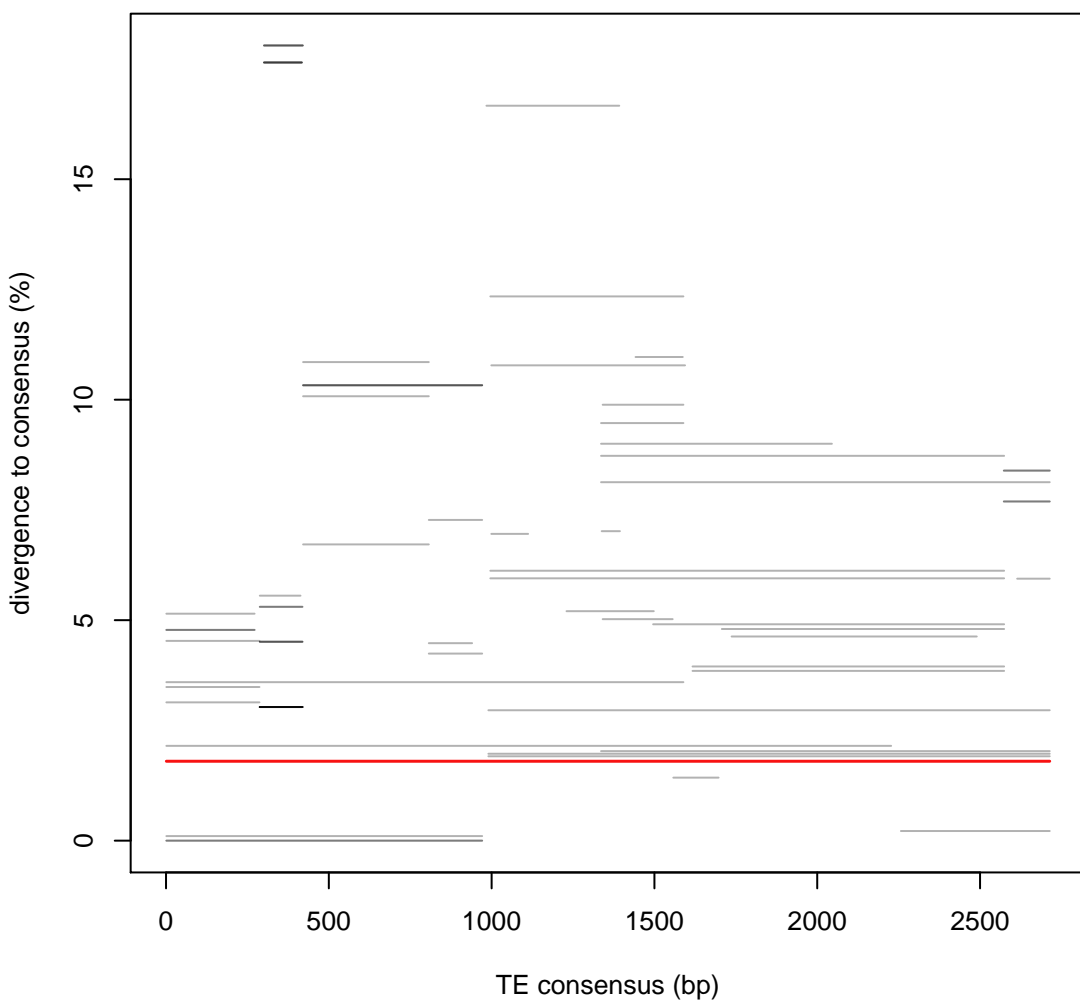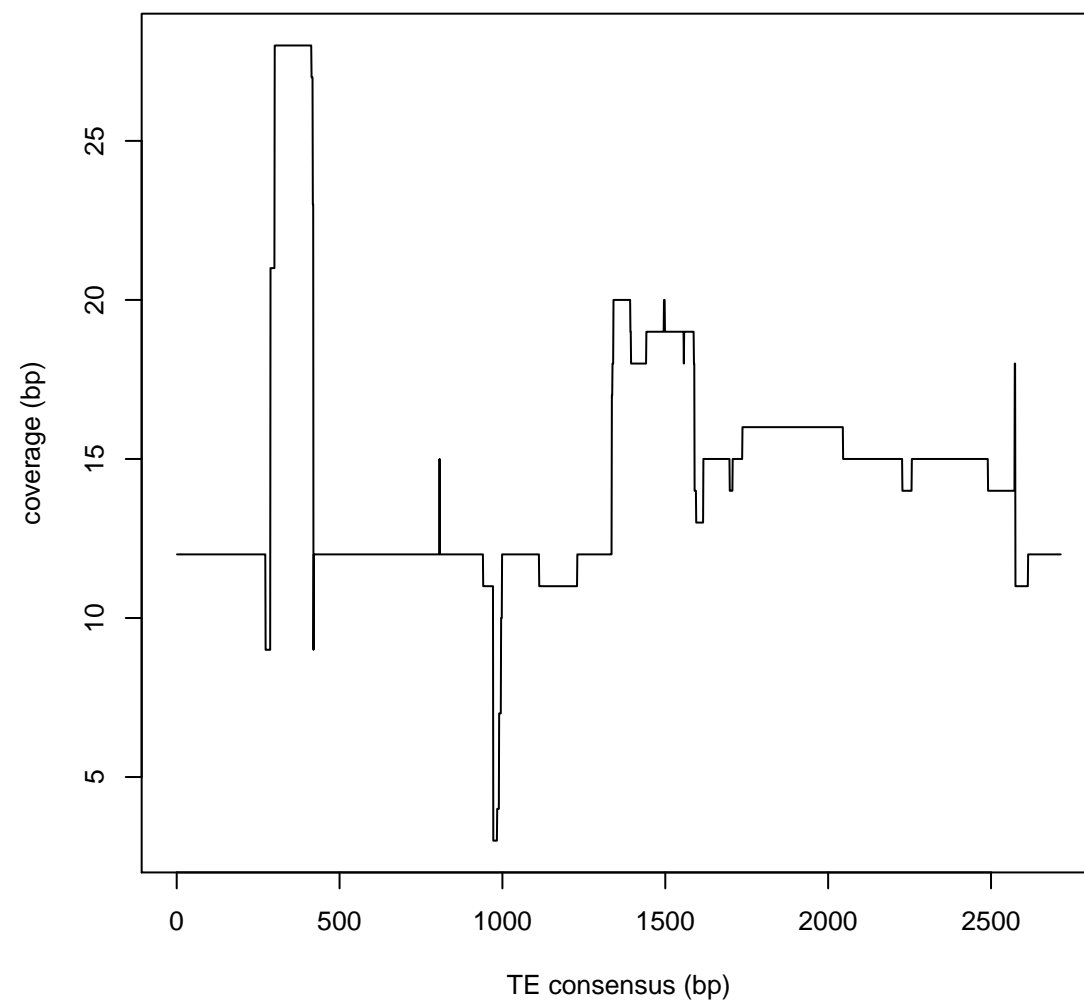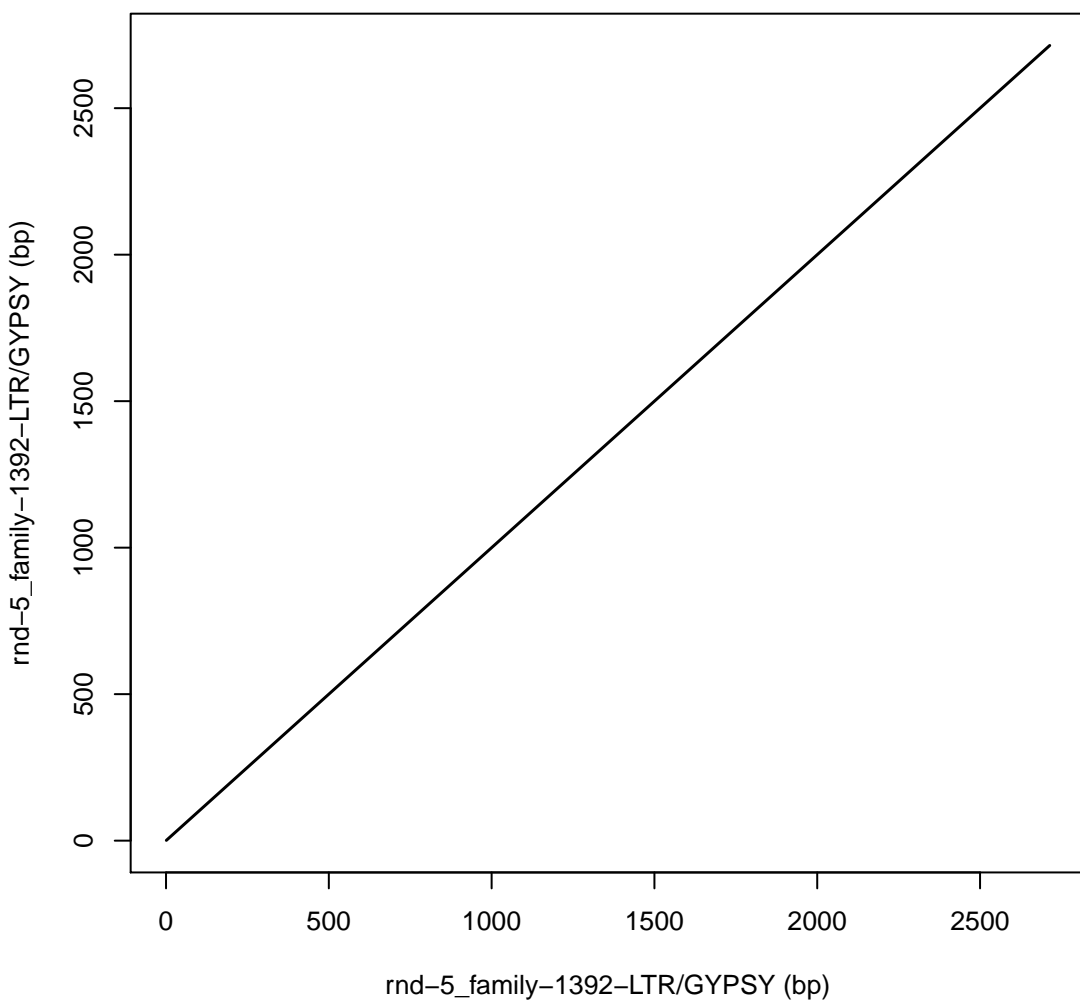**consensus size: 2714bp; fragments: 74; full length: 1 (>=2442.6bp)**

TE consensus (bp)

divergence to consensus (%)

TE consensus genomic coverage

coverage (bp)

TE consensus (bp)

TE consensus self dotplot (blastn)

rnd−5_family−1392−LTR/GYPSY (bp)

rnd−5_family−1392−LTR/GYPSY (bp)

TE consensus structure and protein hits

Gypsy−2_DSe_gag LTR/Gypsy

YOYO_gag LTR/Gypsy

rnd−5_family−1392−LTR/GYPSY (bp)