TE: rnd-1_family-396-LTR/GYPSY
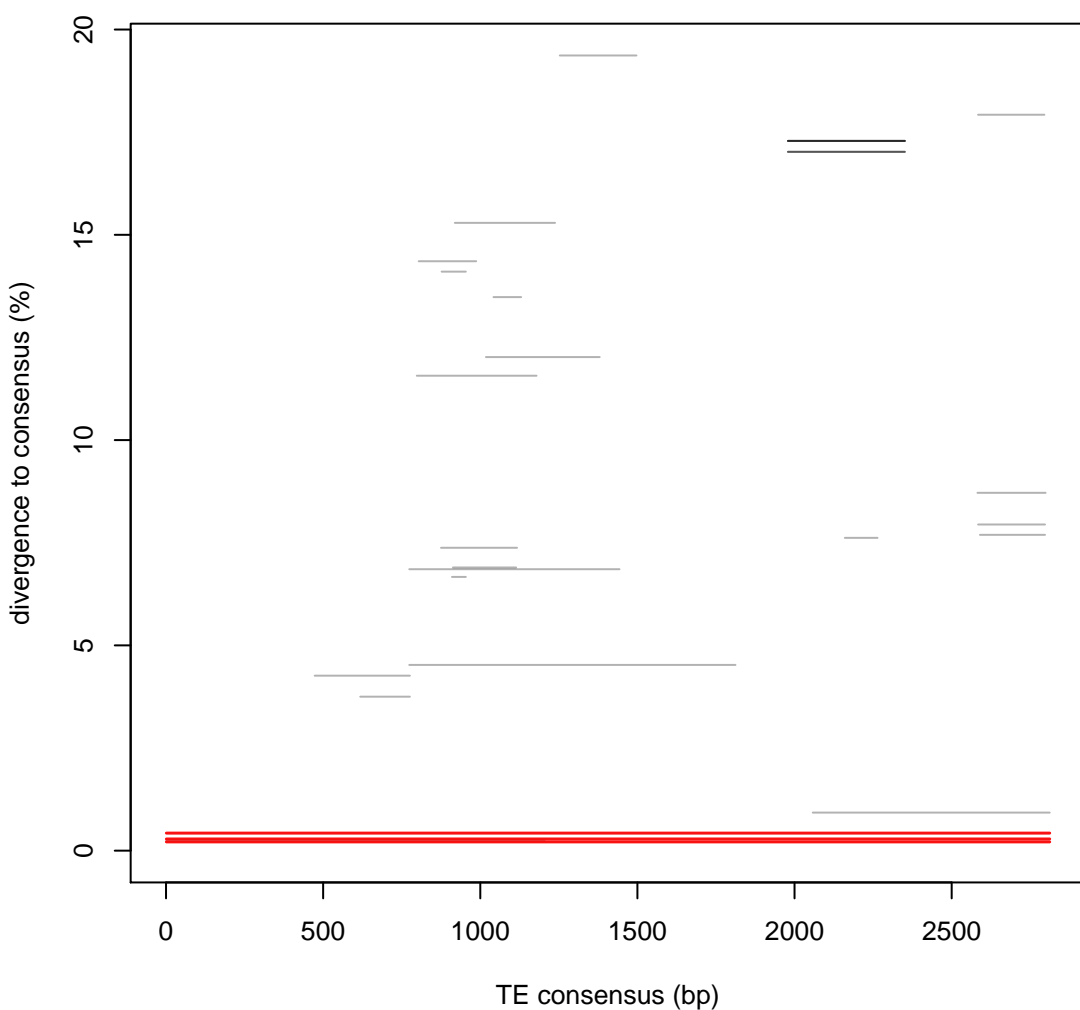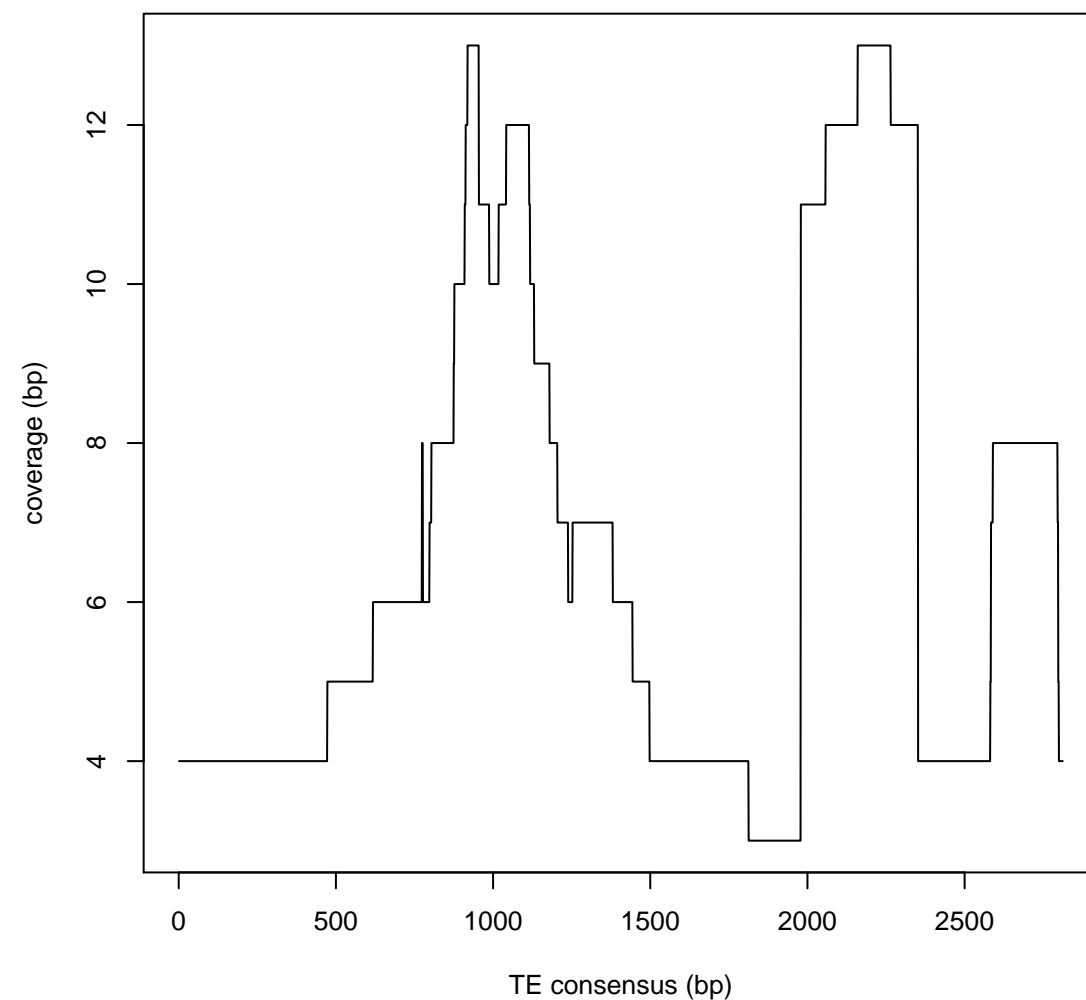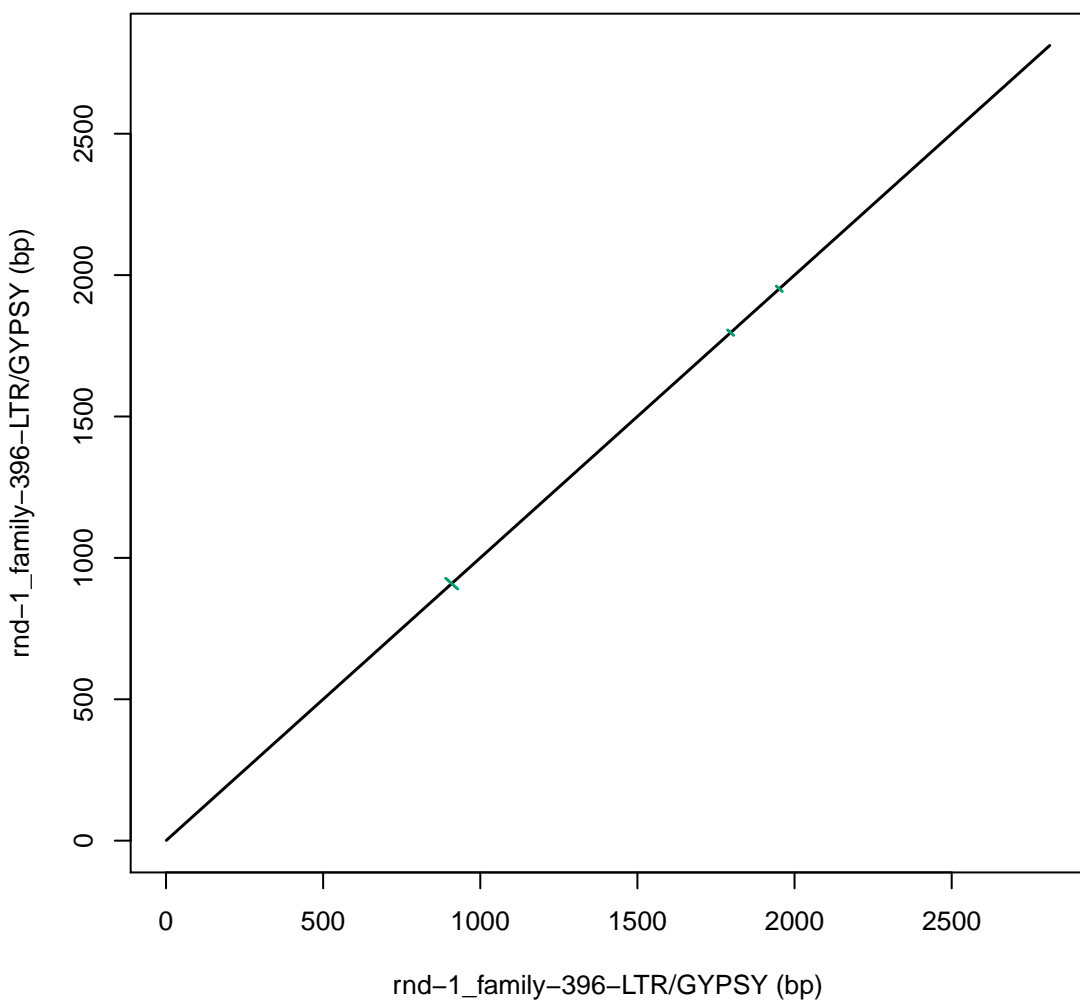**consensus size: 2812bp; fragments: 32; full length: 3 (>=2530.8bp)**

TE consensus (bp)

divergence to consensus (%)

TE consensus genomic coverage

coverage (bp)

TE consensus (bp)

TE consensus self dotplot (blastn)

rnd-1_family-396-LTR/GYPSY (bp)

rnd-1_family-396-LTR/GYPSY (bp)

TE consensus structure and protein hits

Gypsy-44_DWil_env LTR/Gypsy

rnd-1_family-396-LTR/GYPSY (bp)