TE: rnd–1_family–271_s_1–LINE/I–Jockey
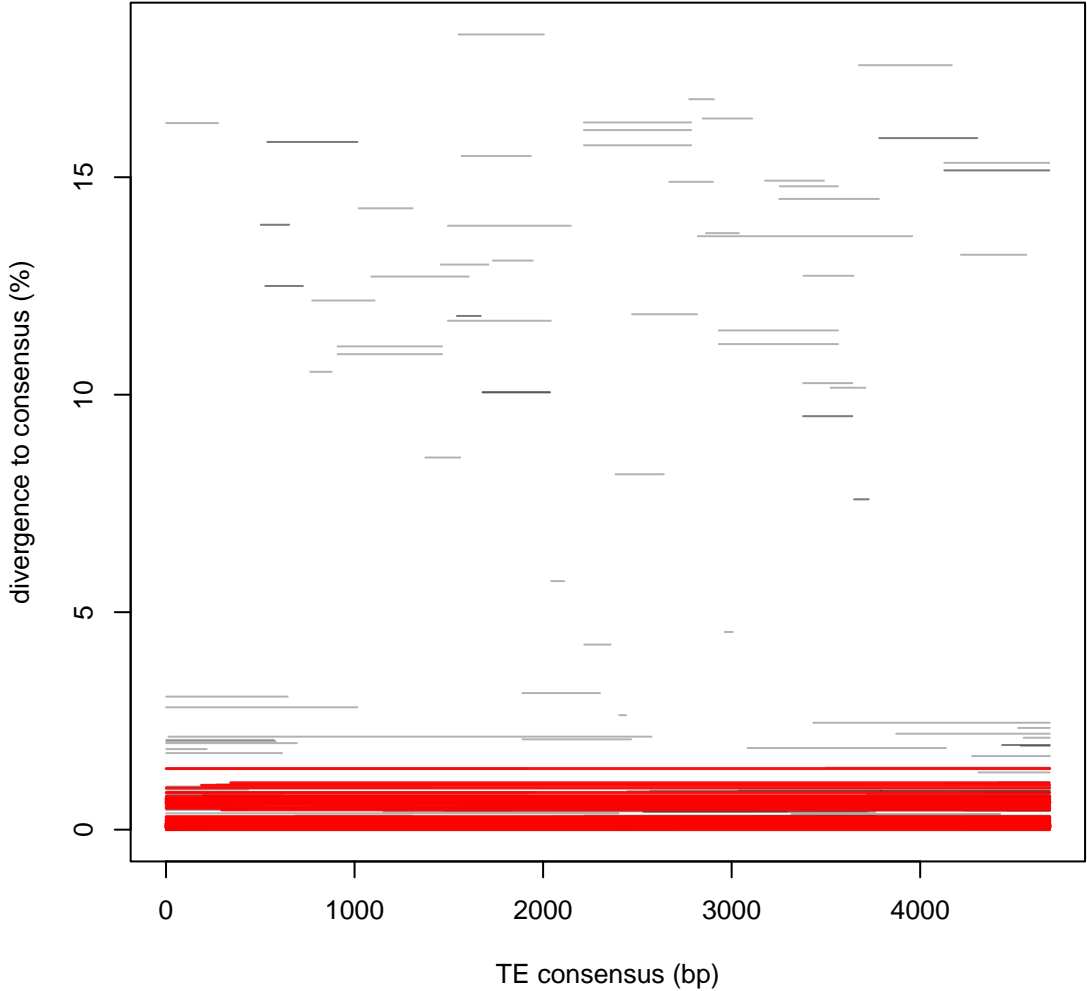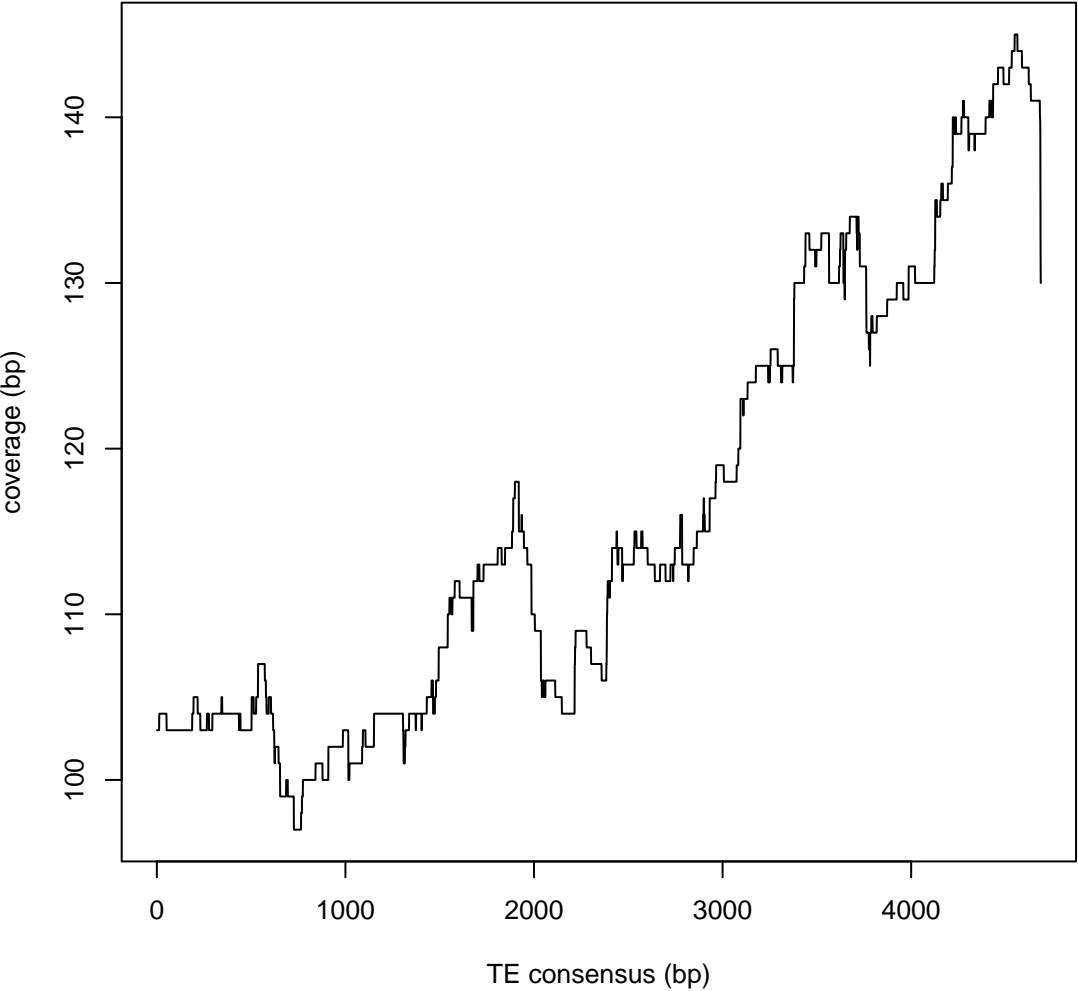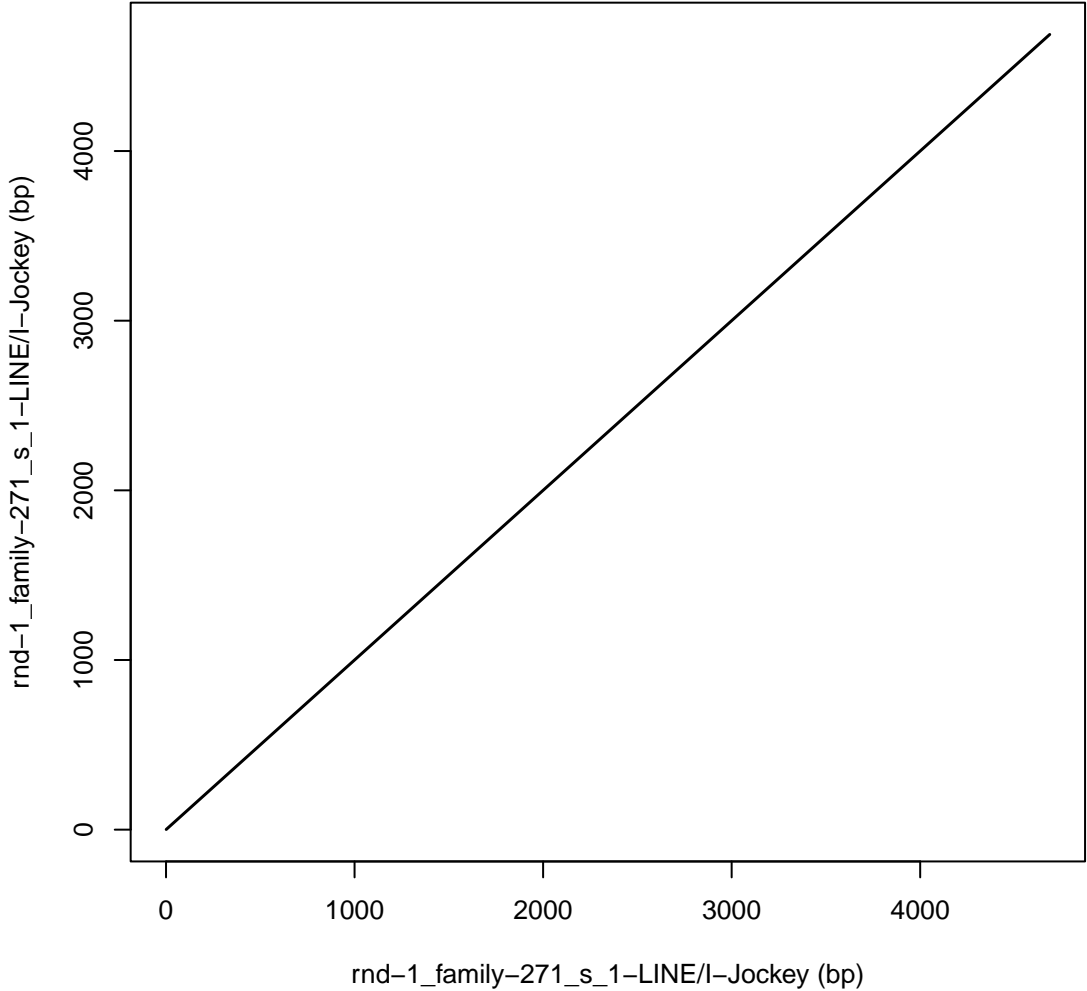consensus size: 4687bp; fragments: 262; full length: 75 (>=4218.3bp)

divergence to consensus (%)

TE consensus (bp)

TE consensus genomic coverage

coverage (bp)

TE consensus (bp)

TE consensus self dotplot (blastn)

rnd–1_family–271_s_1–LINE/I–Jockey (bp)

rnd–1_family–271_s_1–LINE/I–Jockey (bp)

TE consensus structure and protein hits

psy–17_ST LTR/Gypsy

Gypsy–8_Alp_2p LTR/Gypsy

R1–1_NVi_pol LINE/R1

Transib–2_DAn TIR/CM

rnd–1_family–271_s_1–LINE/I–Jockey (bp)