TE: rnd−1_family−77_s_0−LTR/Gypsy
consensus size: 7152bp; fragments: 913; full length: 9 (>=6436.8bp)

divergence to consensus (%)
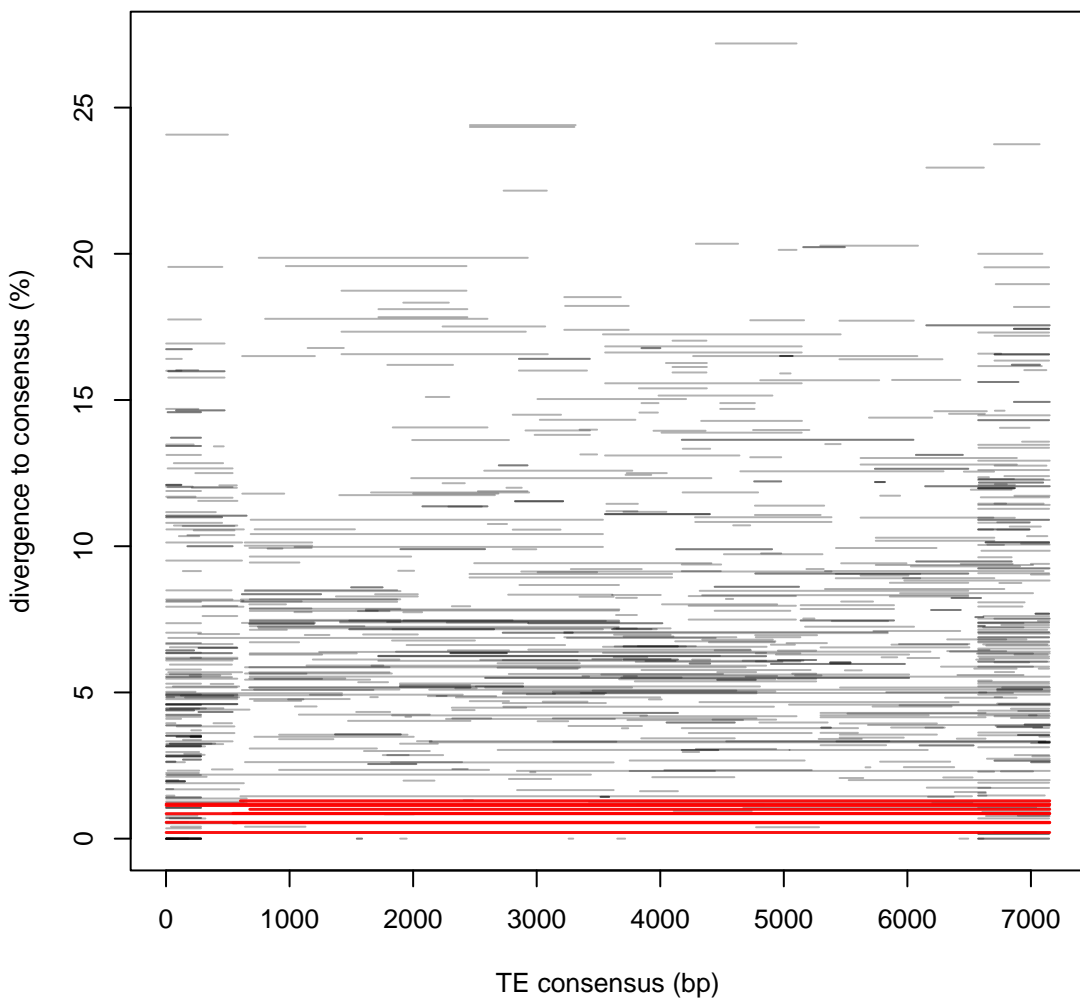
TE consensus (bp)

TE consensus genomic coverage

coverage (bp)
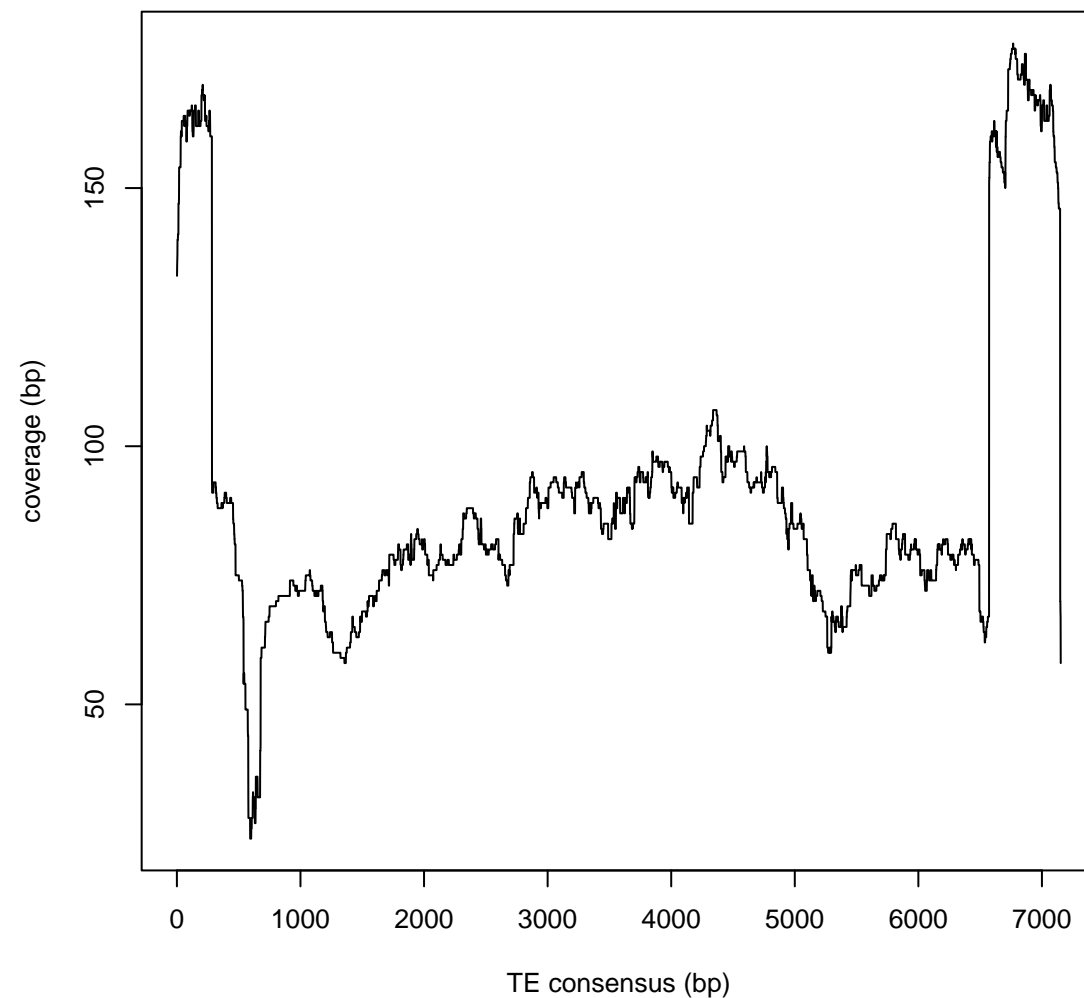
TE consensus (bp)

TE consensus self dotplot (blastn)

rnd−1_family−77_s_0−LTR/Gypsy (bp)

rnd−1_family−77_s_0−LTR/Gypsy (bp)
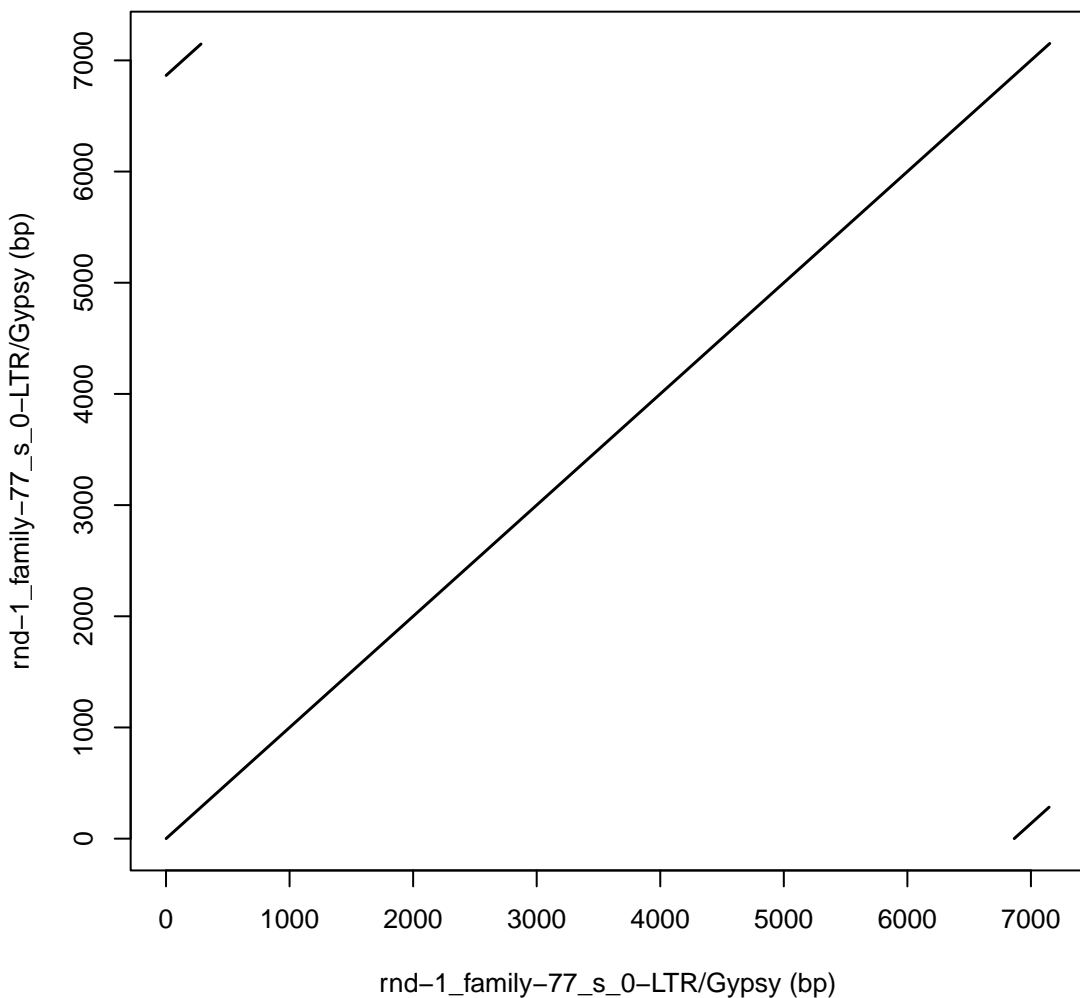
TE consensus structure and protein hits

no orf >400 bp detected

rnd−1_family−77_s_0−LTR/Gypsy (bp)