TE: rnd−5_family−2779−LTR/Pao
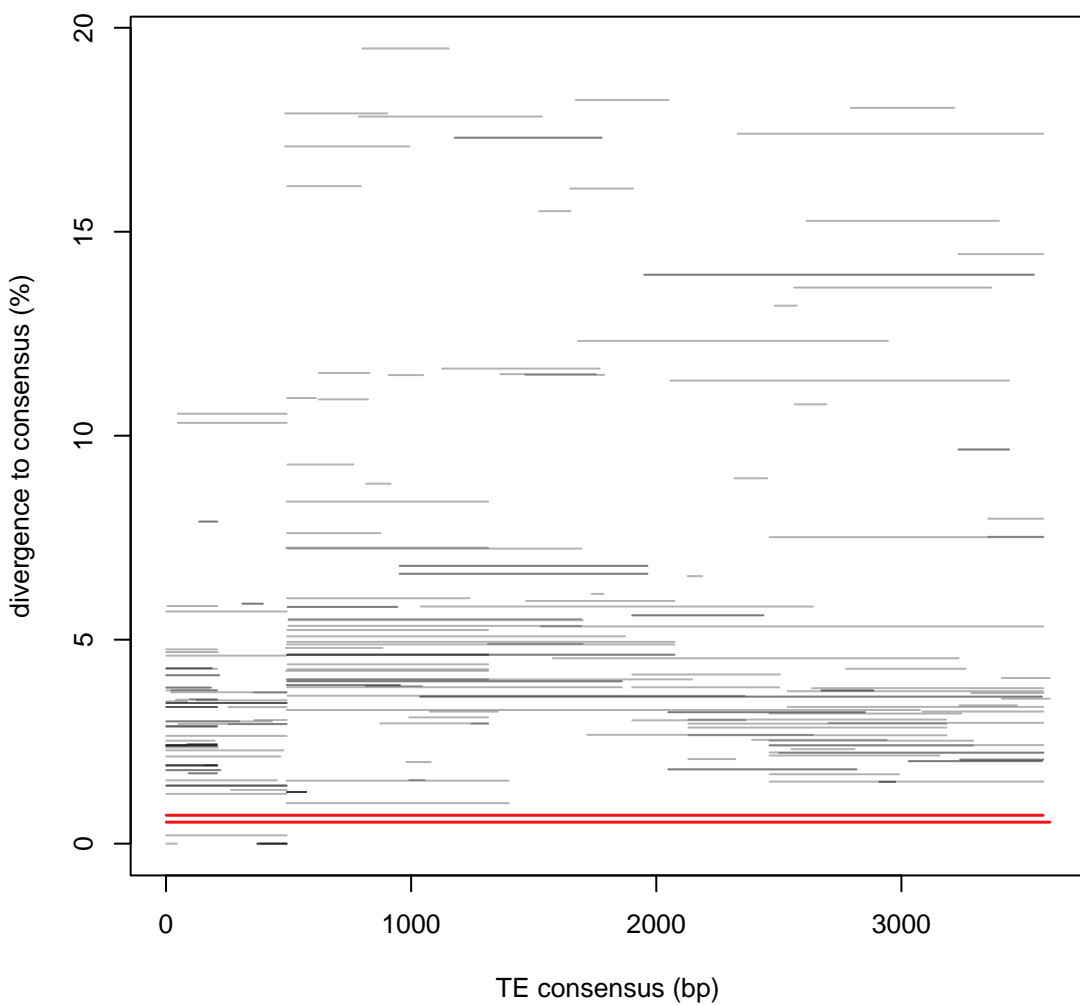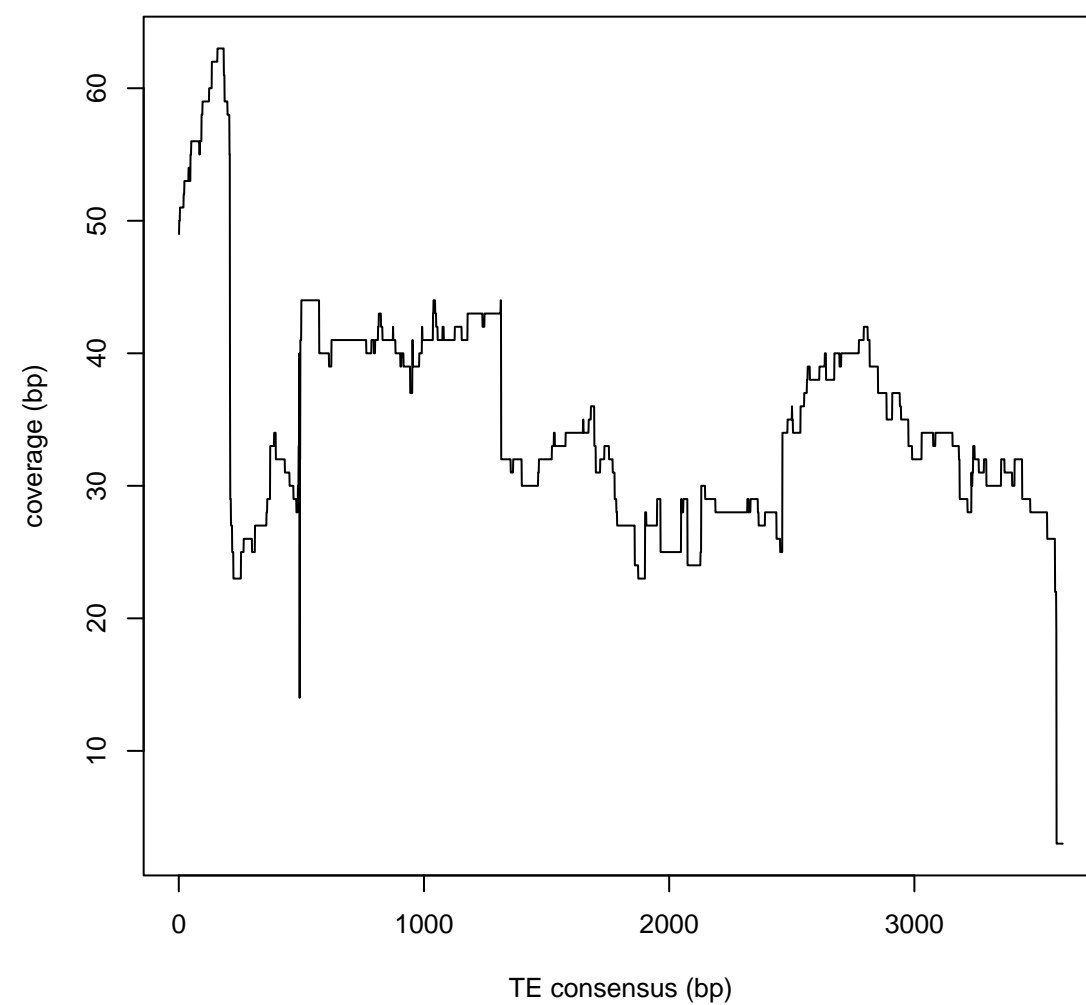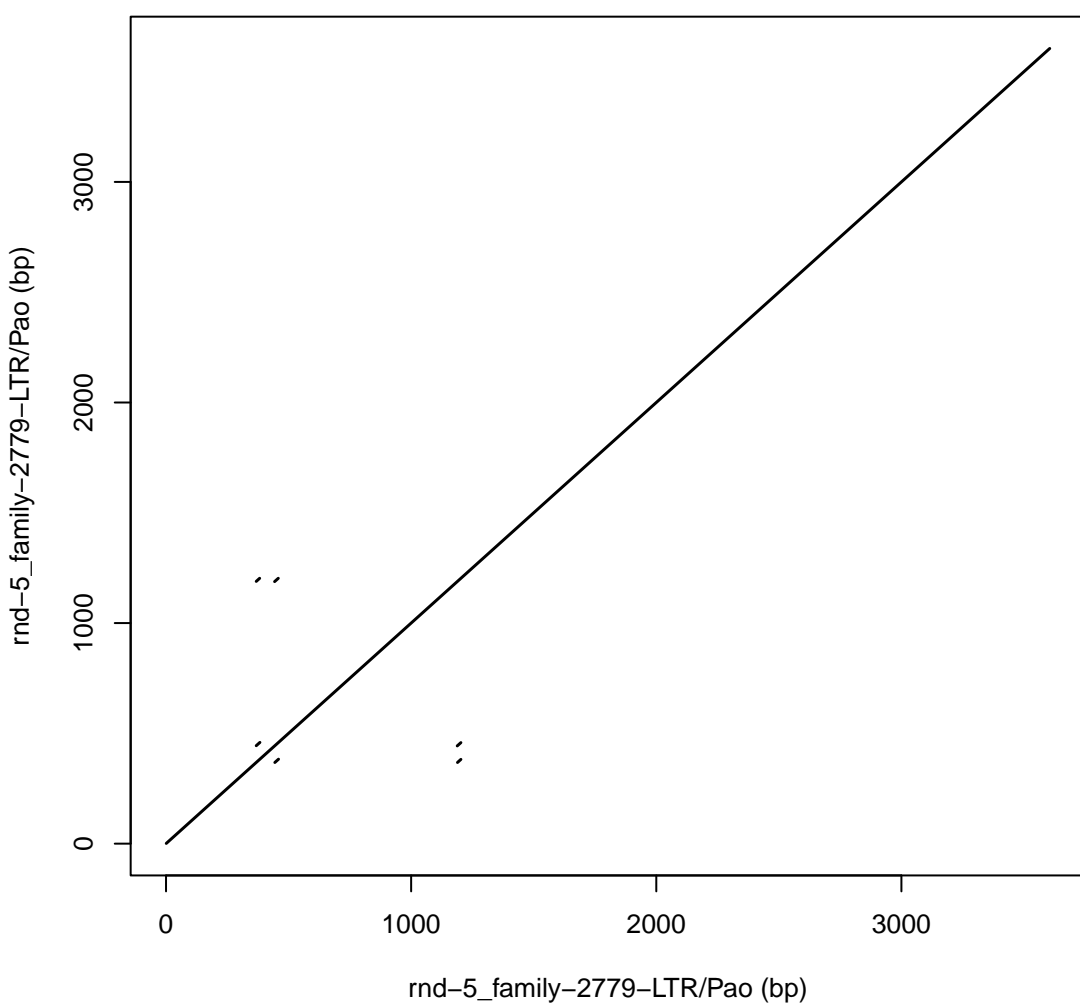consensus size: 3605bp; fragments: 215; full length: 2 (>=3244.5bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

Gypsy45_DR LTR/Gypsy
BEL−1_Adi_pol LTR/Pao
NINJA_pol LTR/Pao
BEL29_DR_pol LTR/Pa