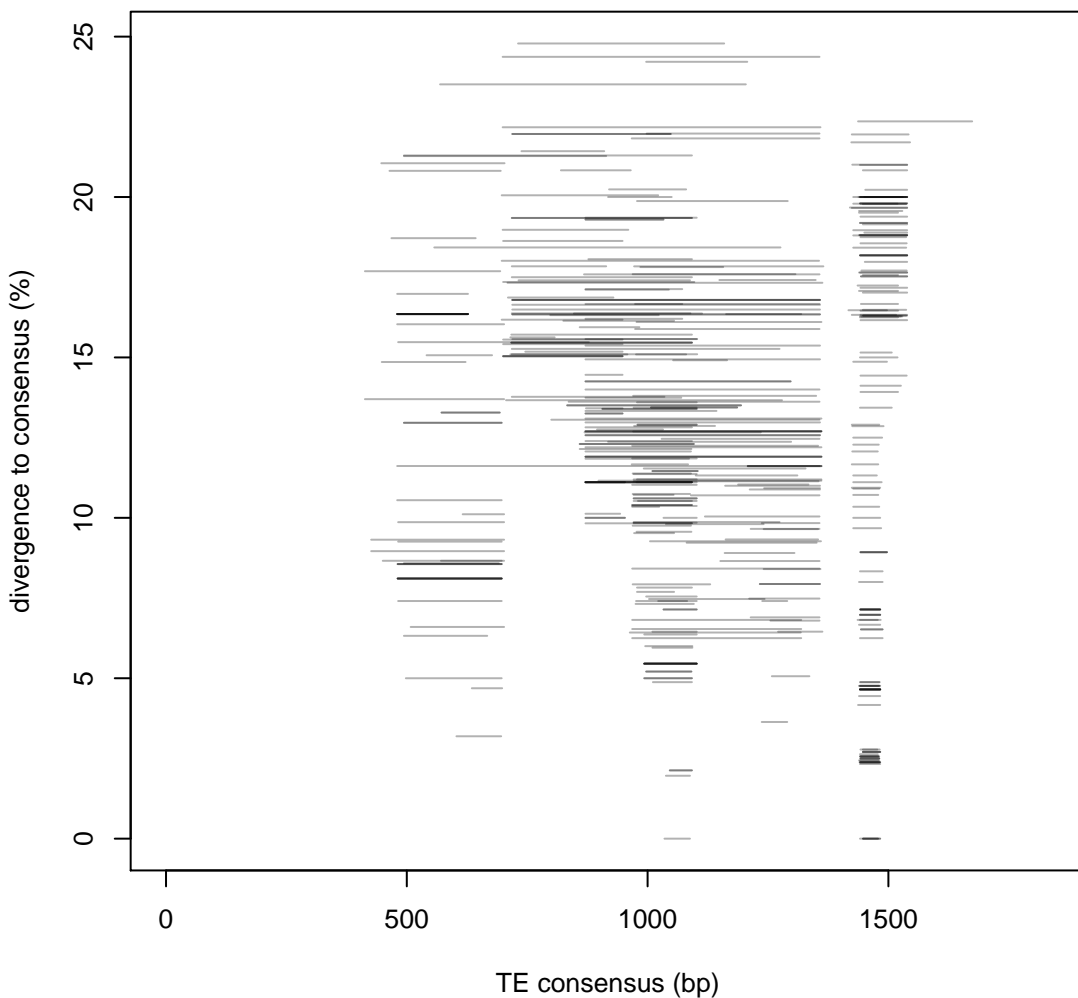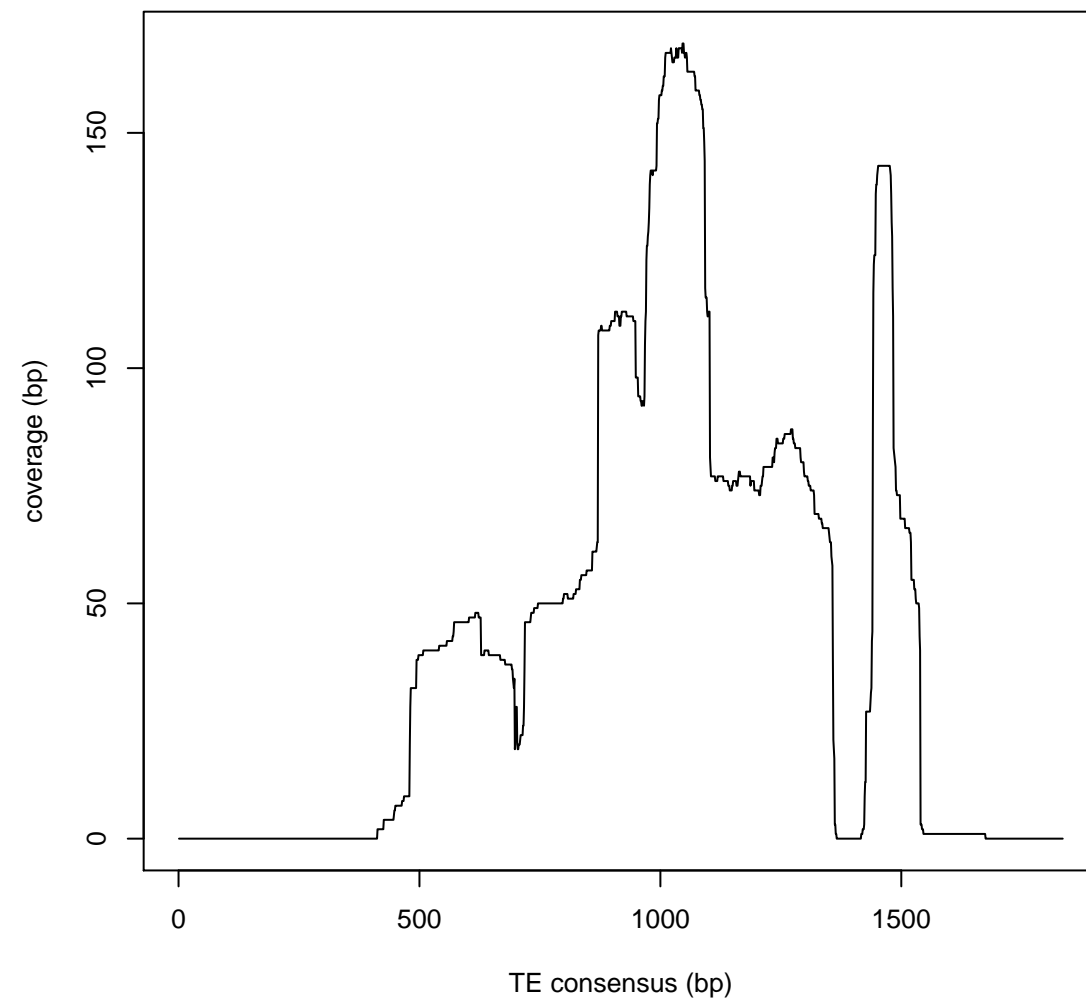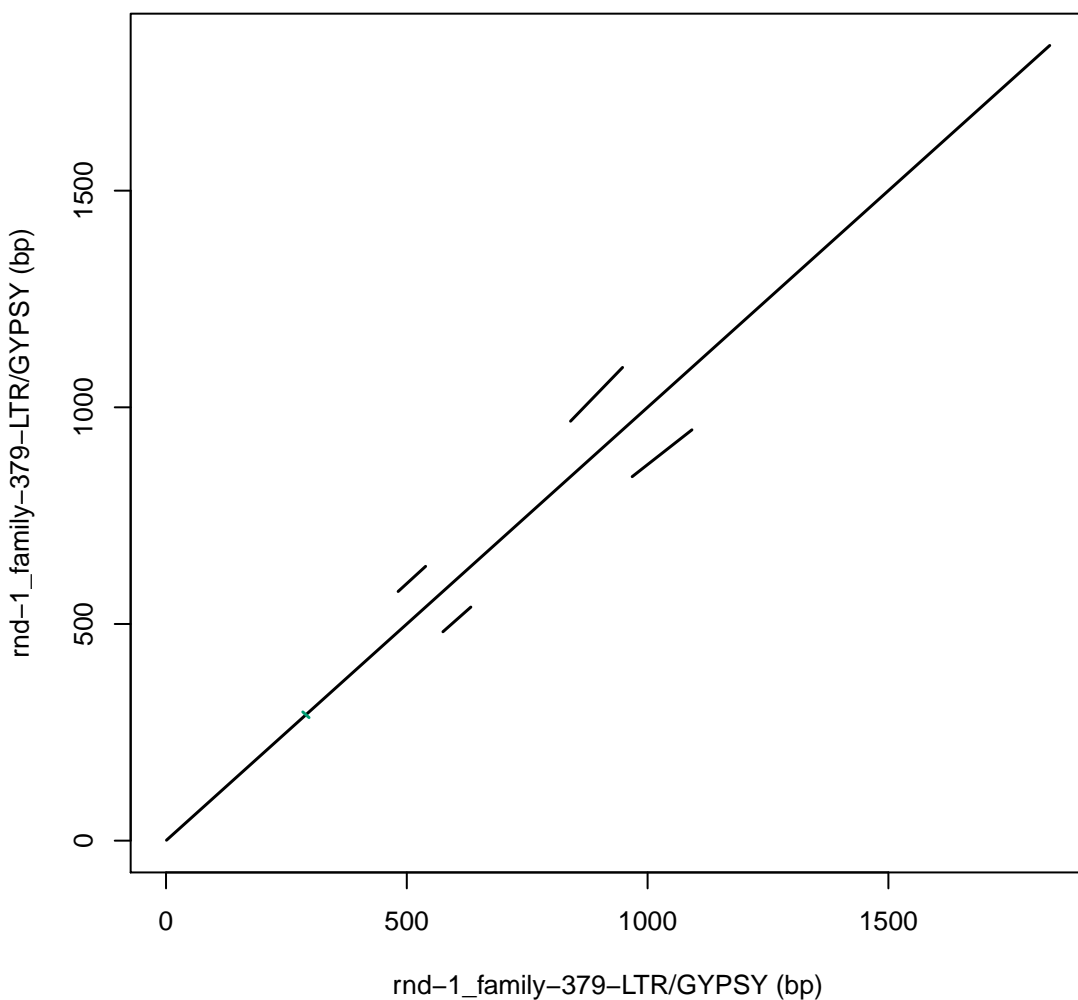**TE: rnd−1_family−379−LTR/GYPSY**
consensus size: 1835bp; fragments: 420; full length: 0 (>=1651.5bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

MuDR−10_ALy_p1 TIR/MULE−MuDR

RTEX−9_RPr_gag LINE/RTE−X