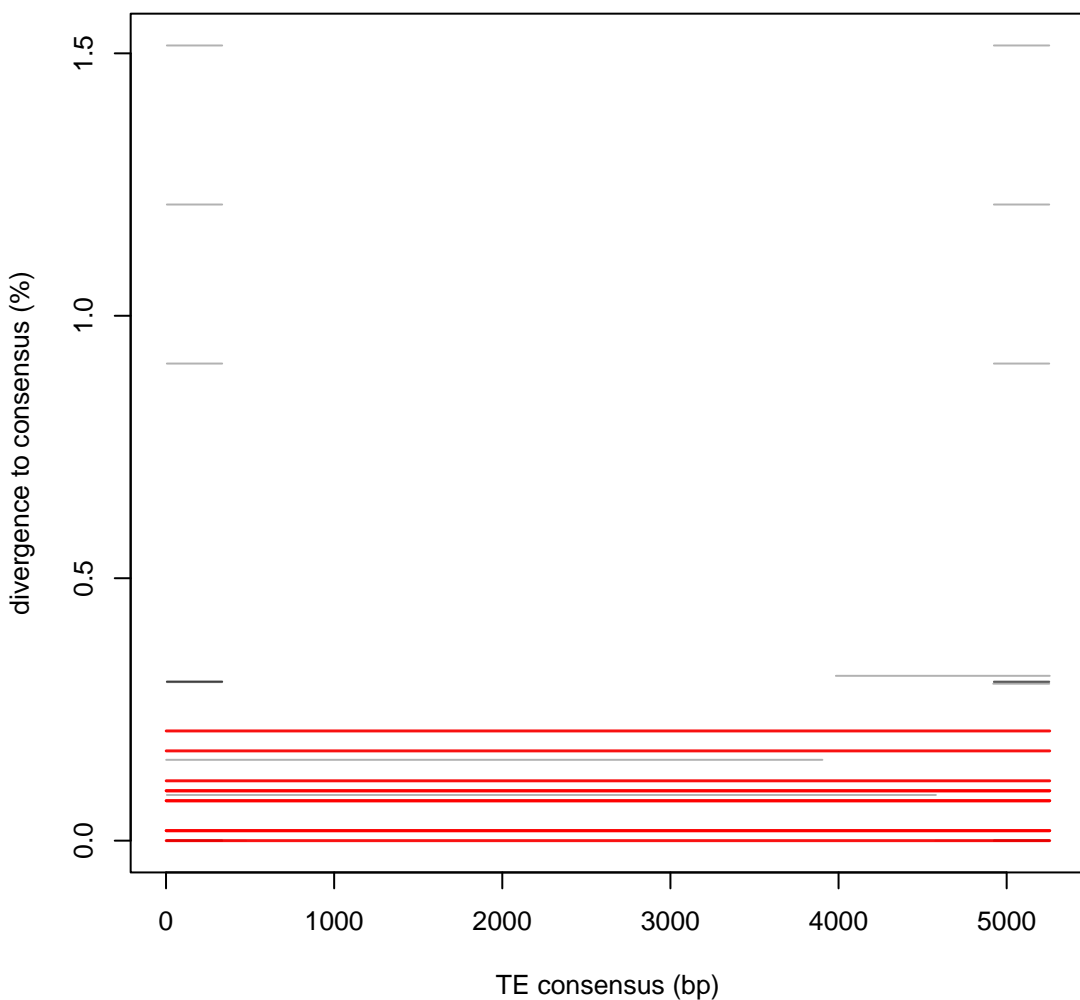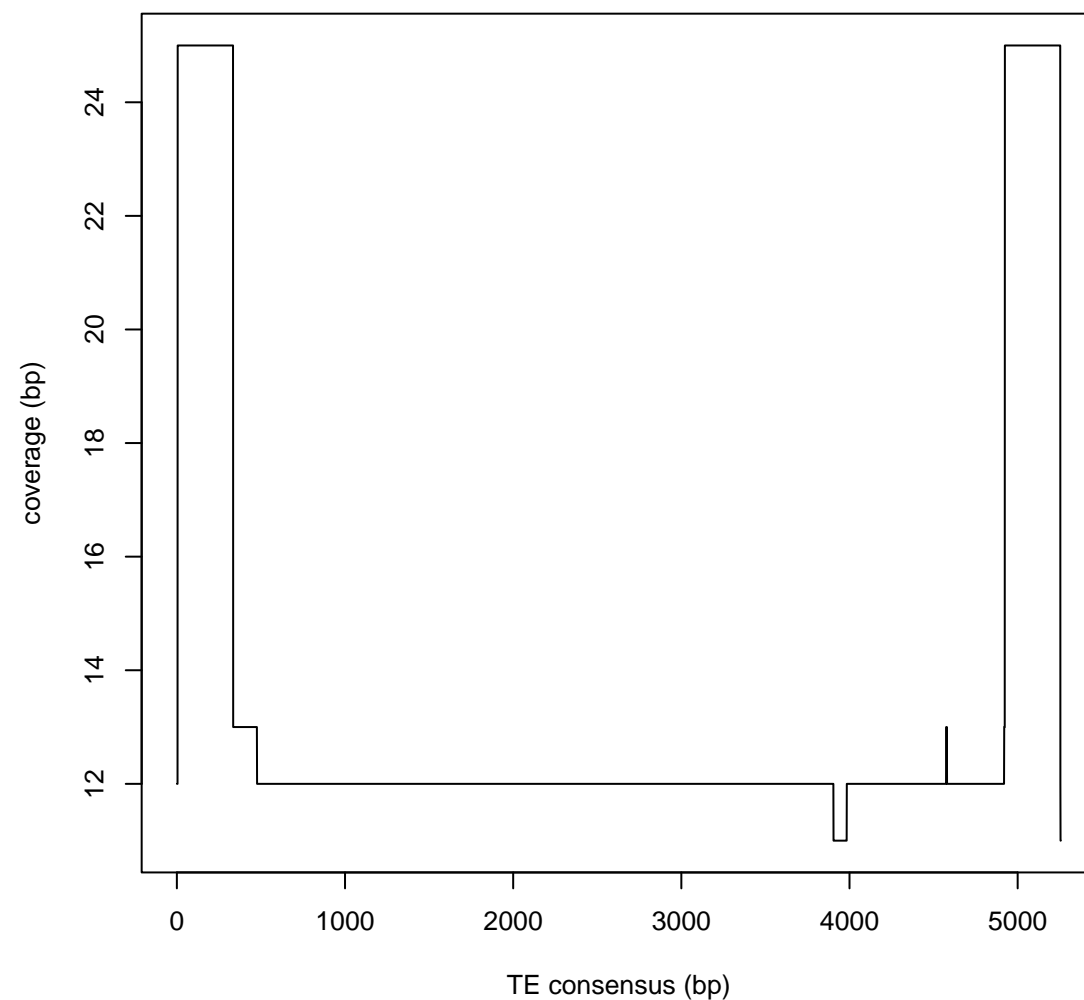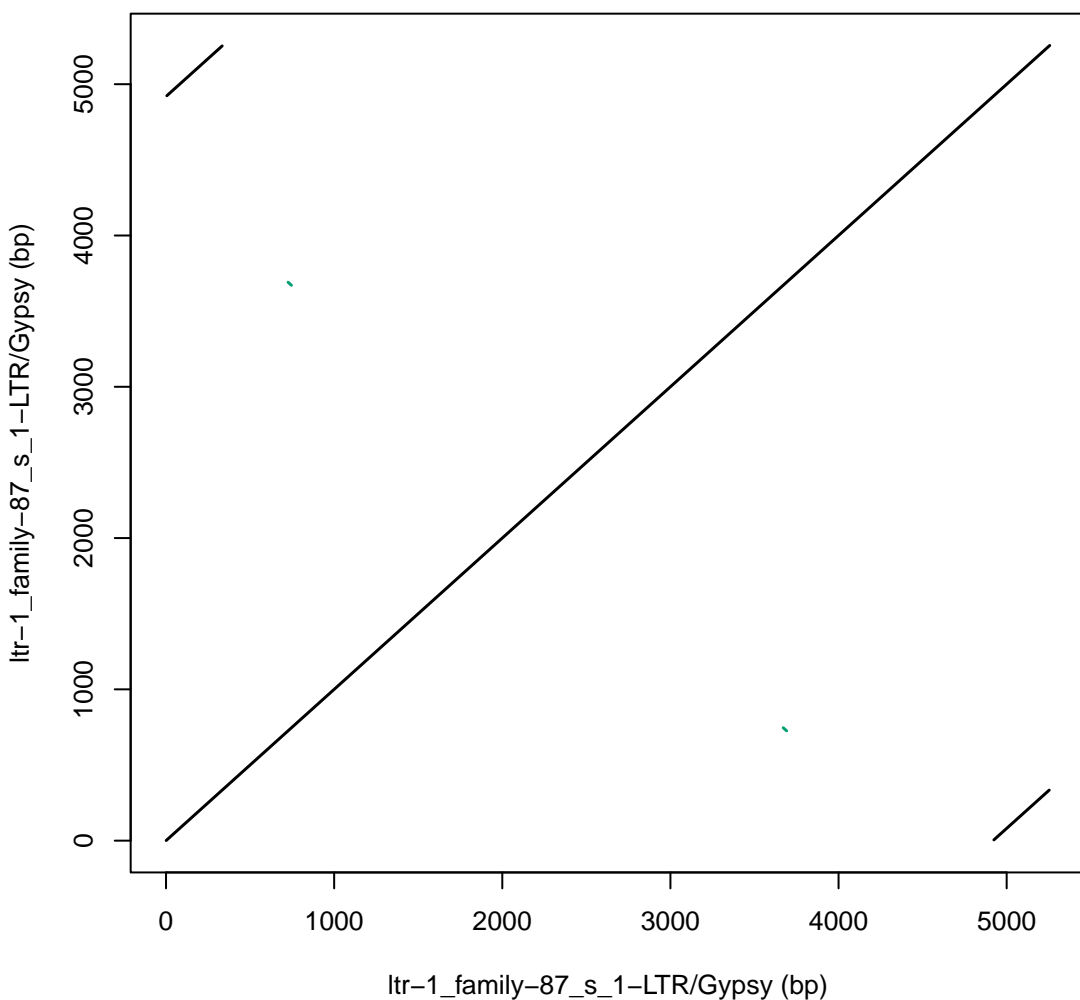**TE: ltr−1_family−87_s_1−LTR/Gypsy**
consensus size: 5256bp; fragments: 40; full length: 10 (>=4730.4bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

TRANSPAC_gag LTR/Gypsy

TRANSPAC_pol LTR/Gypsy