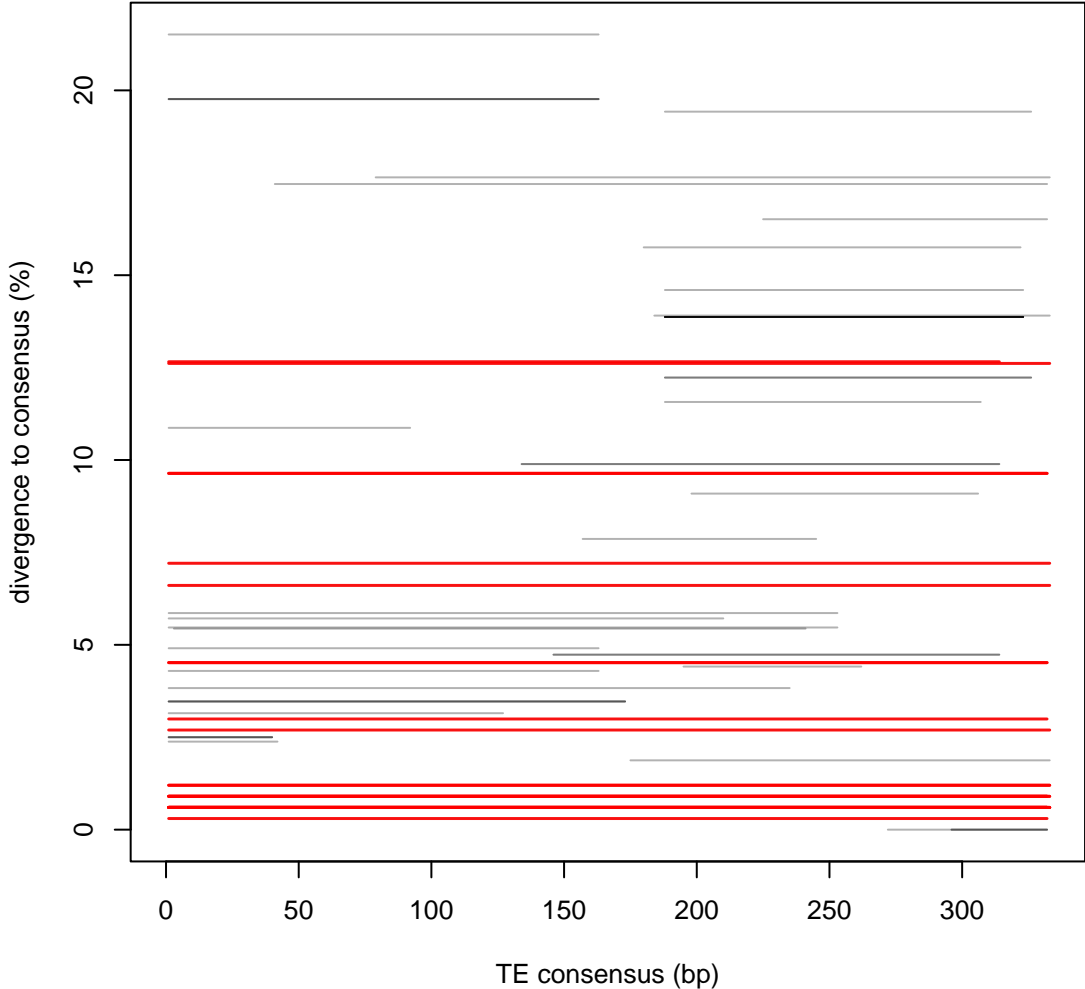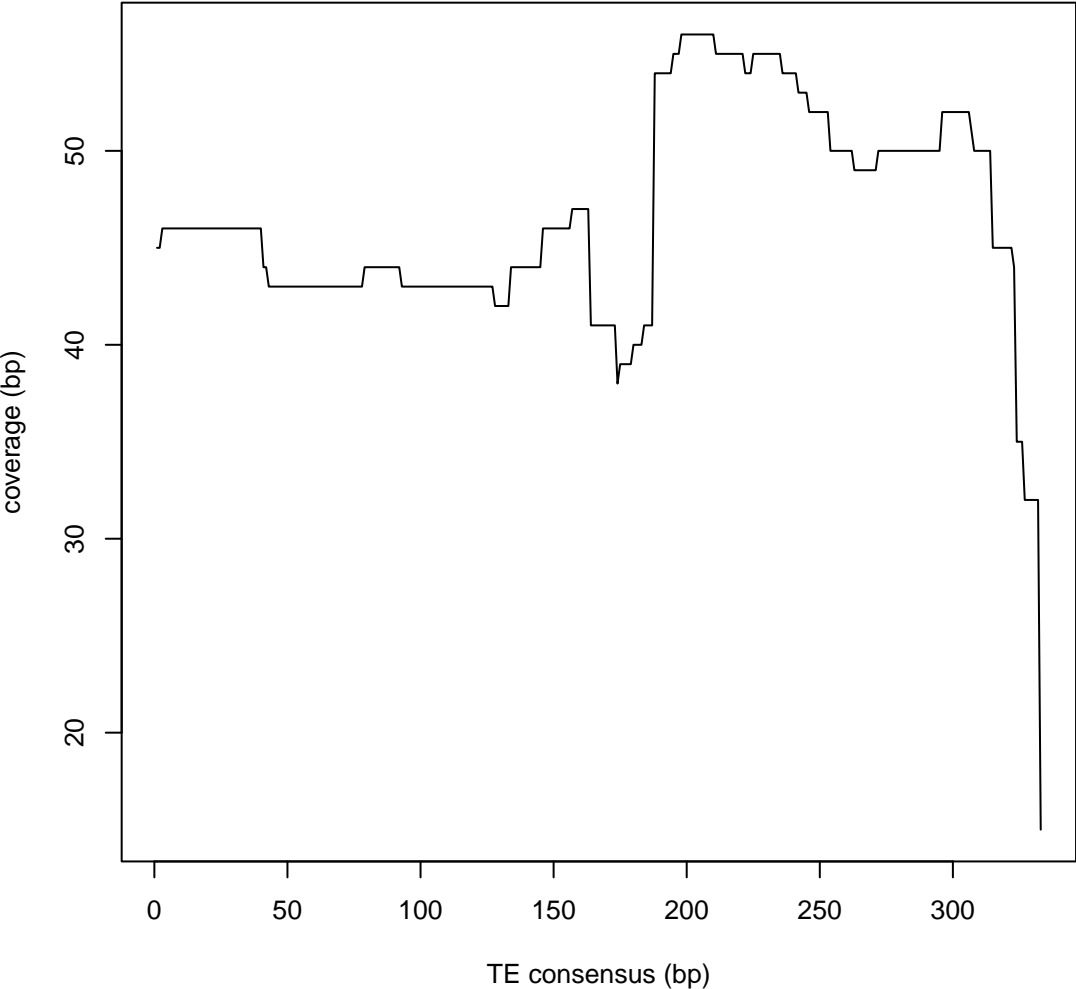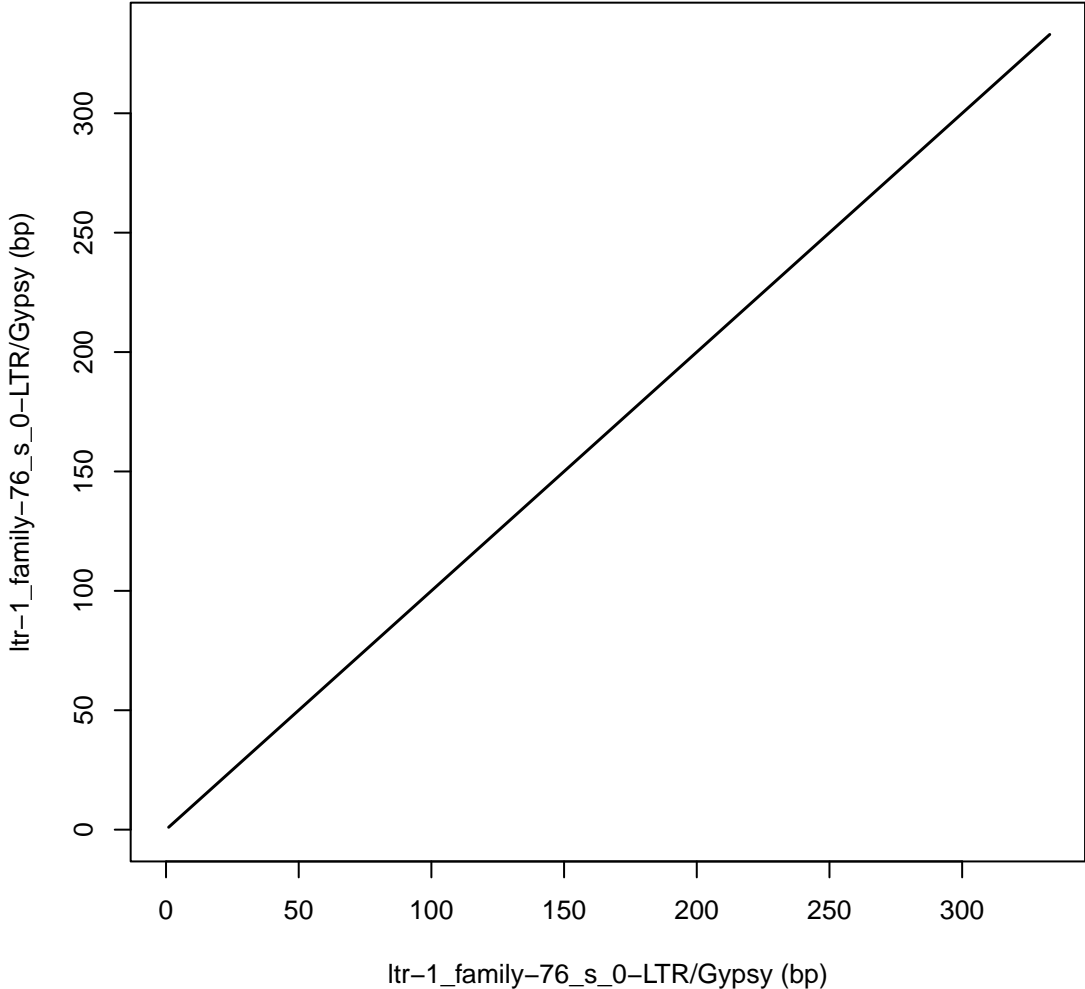**TE: ltr−1_family−76_s_0−LTR/Gypsy**
consensus size: 333bp; fragments: 75; full length: 25 (>=299.7bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected