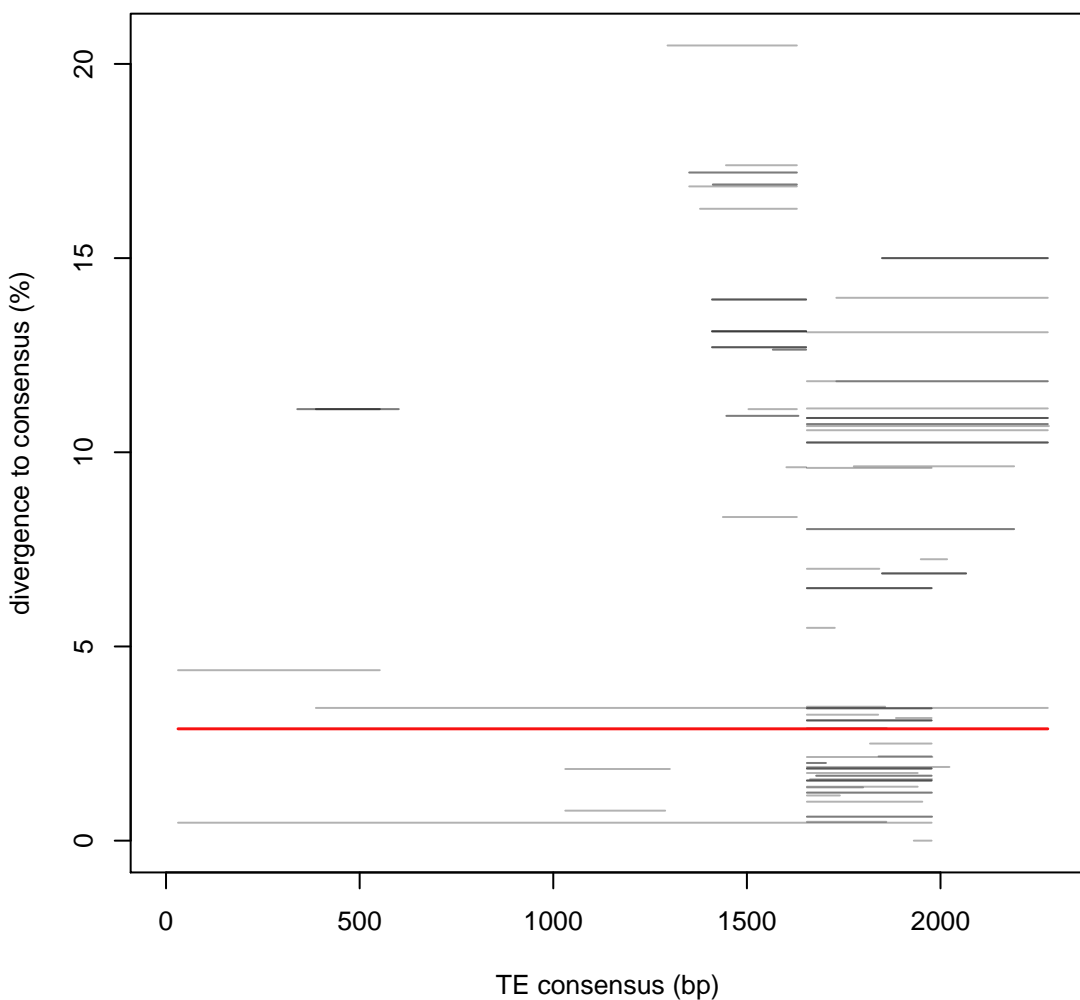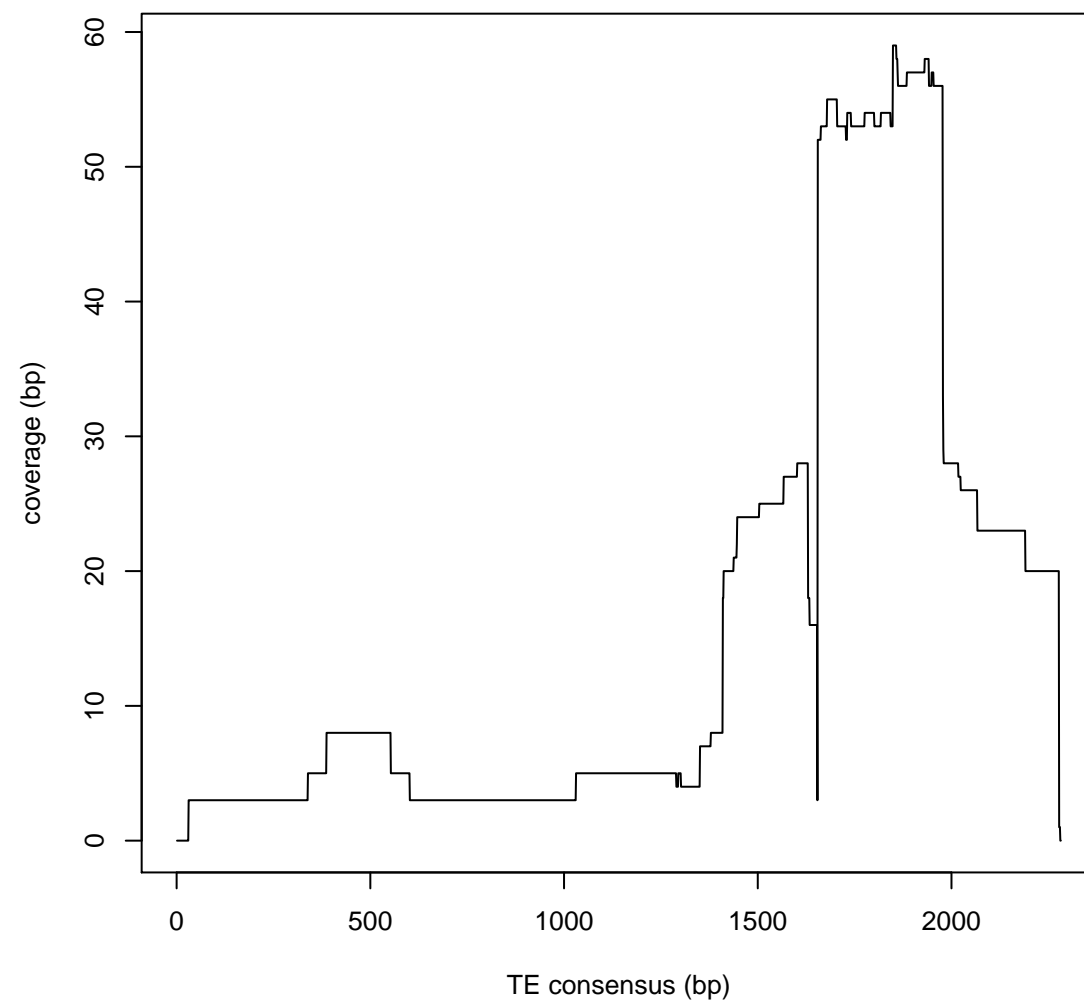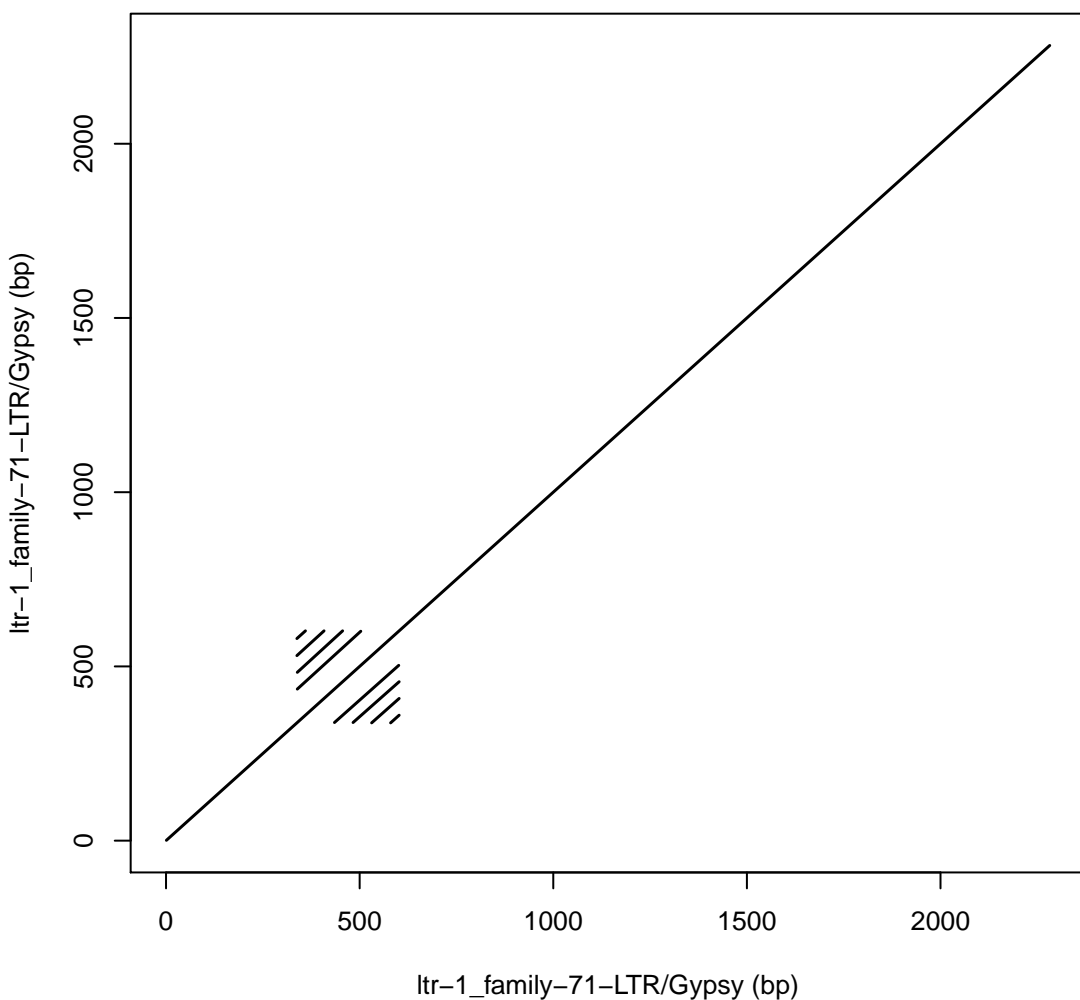**TE: ltr−1_family−71−LTR/Gypsy**
consensus size: 2282bp; fragments: 101; full length: 1 (>=2053.8bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected