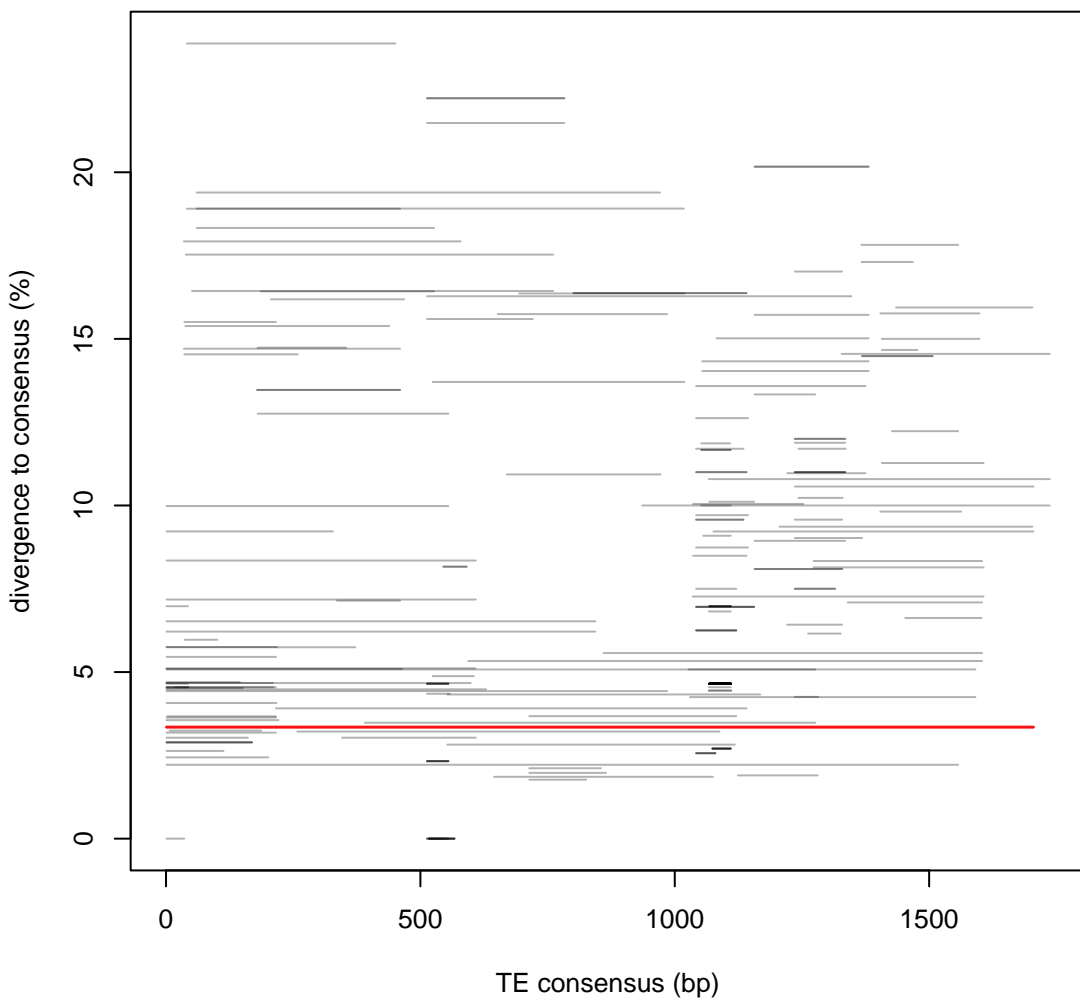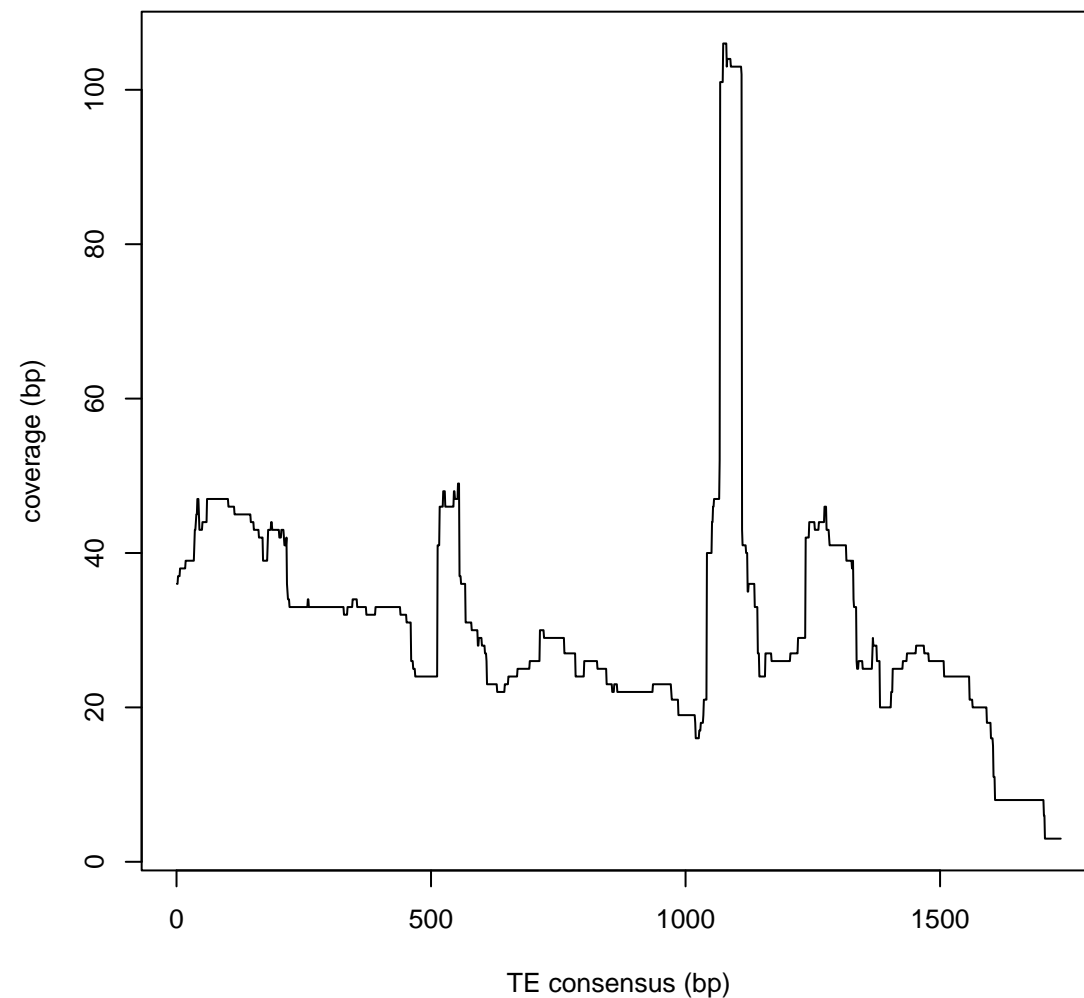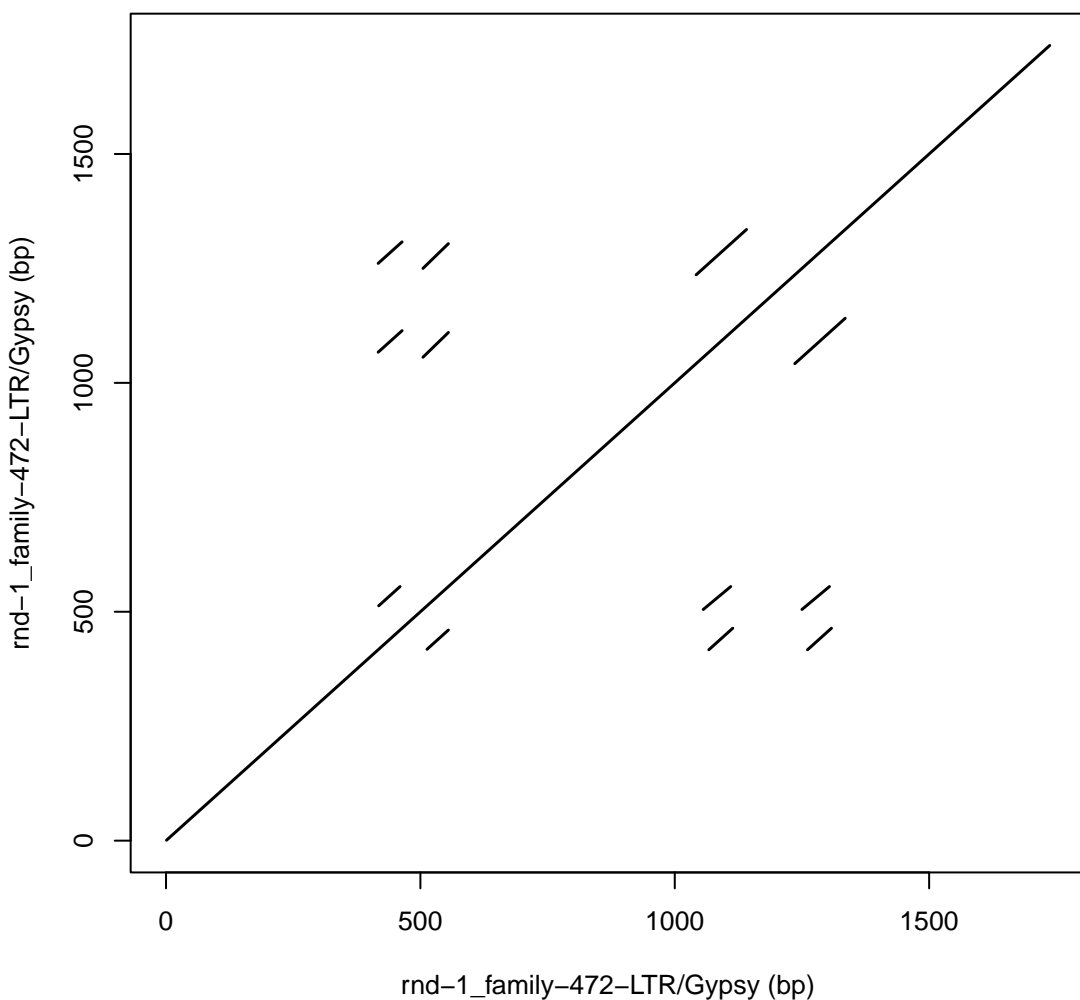**TE: rnd−1_family−472−LTR/Gypsy**
consensus size: 1737bp; fragments: 239; full length: 1 (>=1563.3bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected