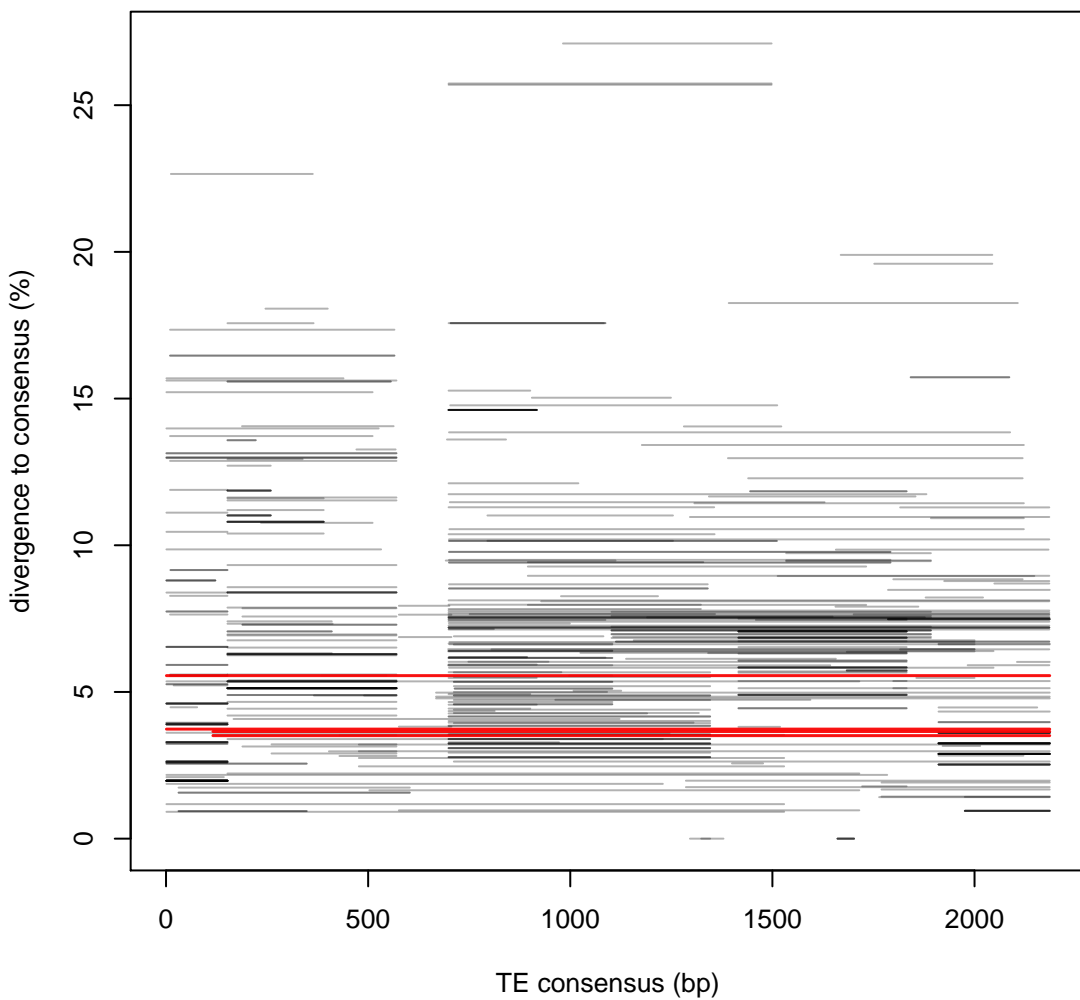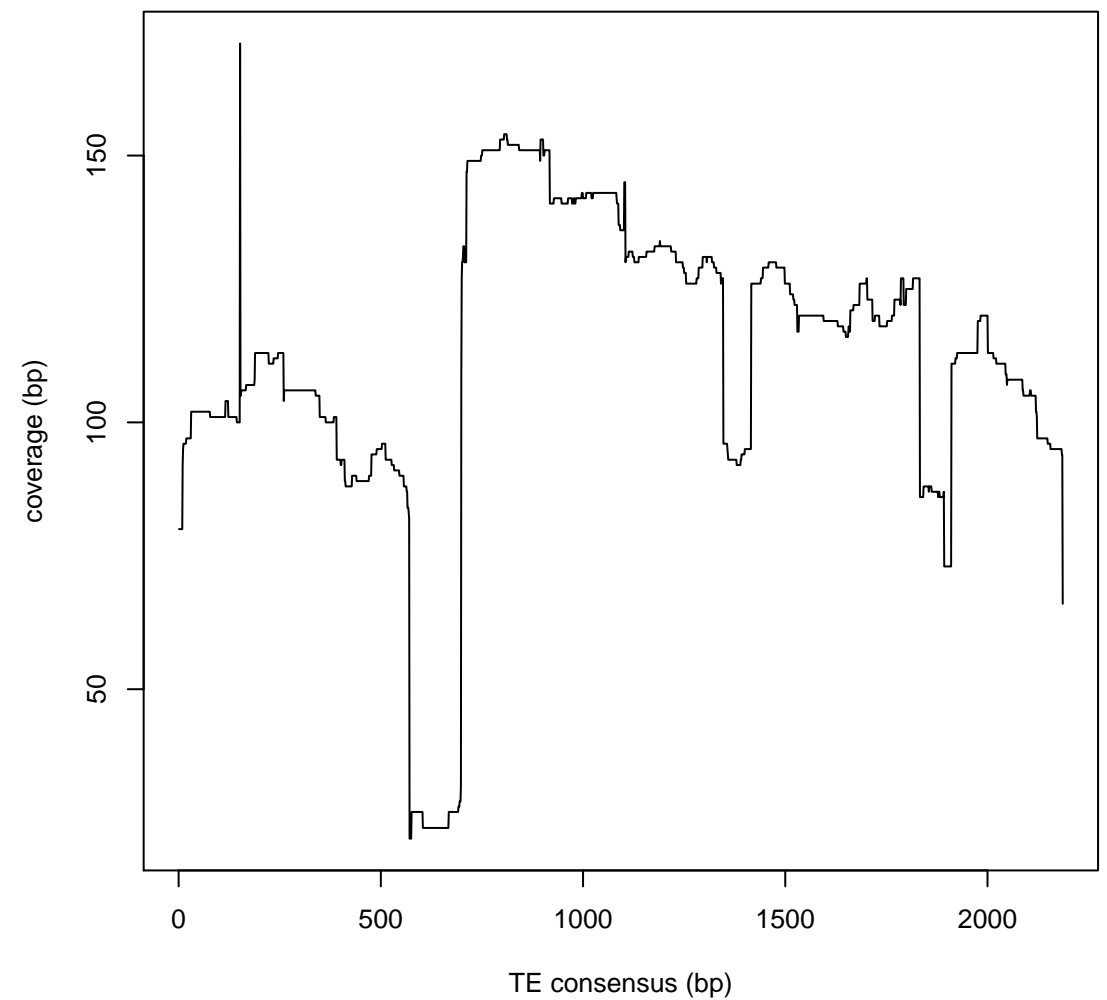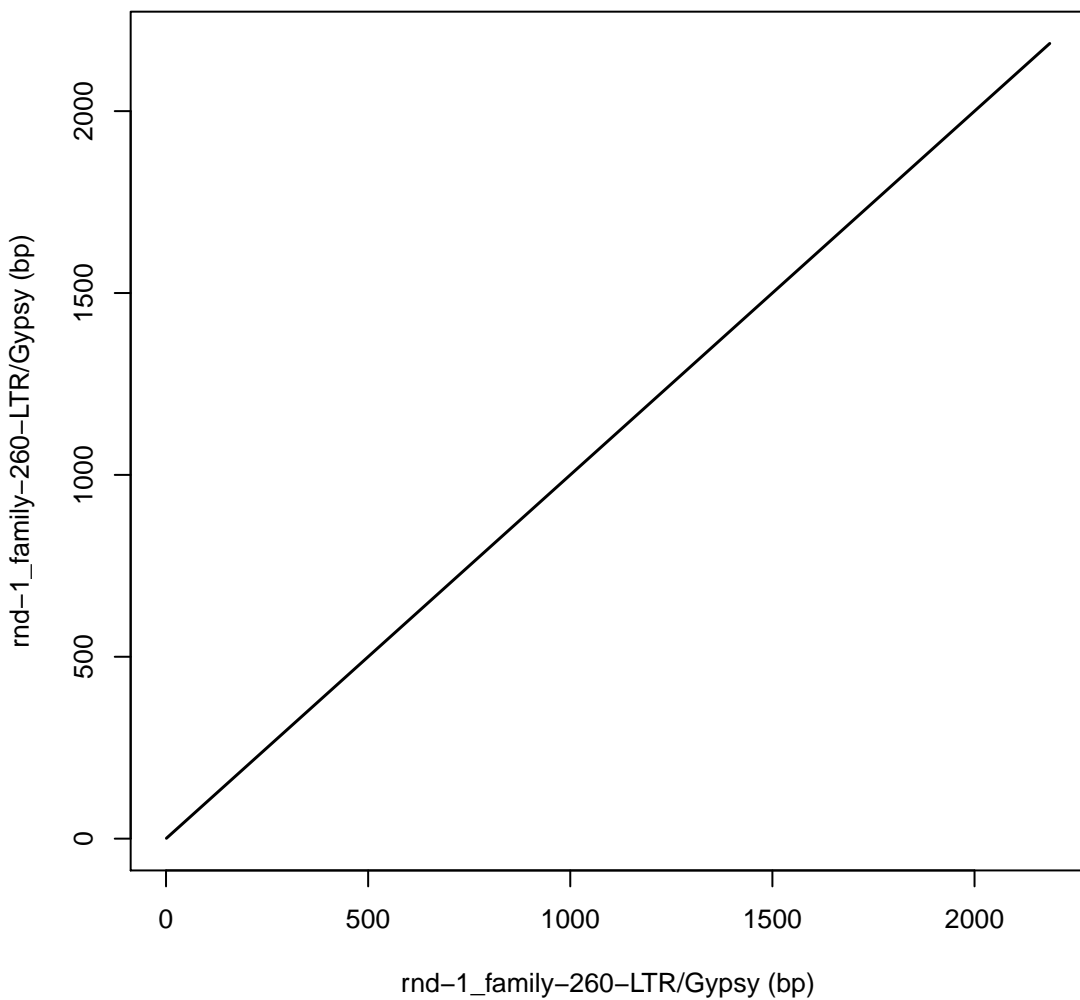**TE: rnd-1_family-260-LTR/Gypsy**
consensus size: 2186bp; fragments: 500; full length: 4 (>=1967.4bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected