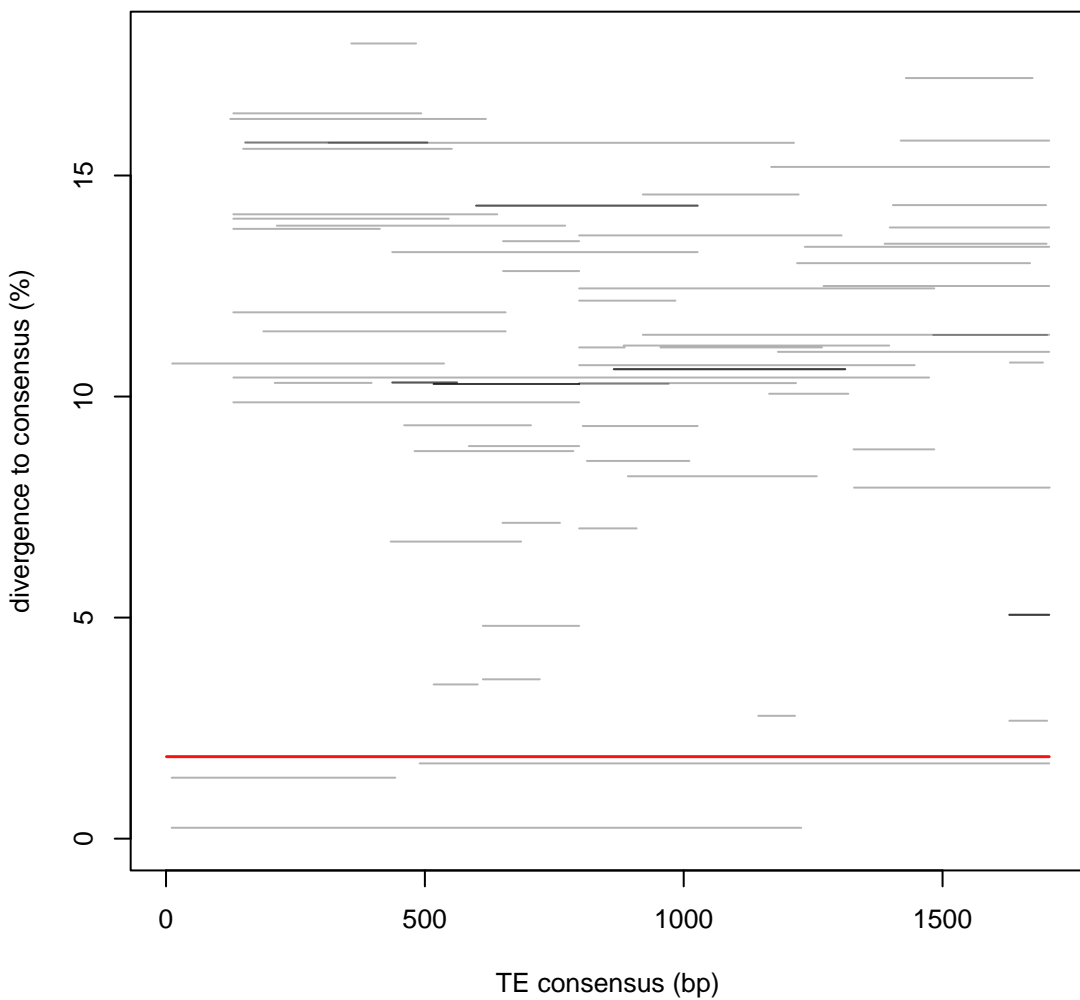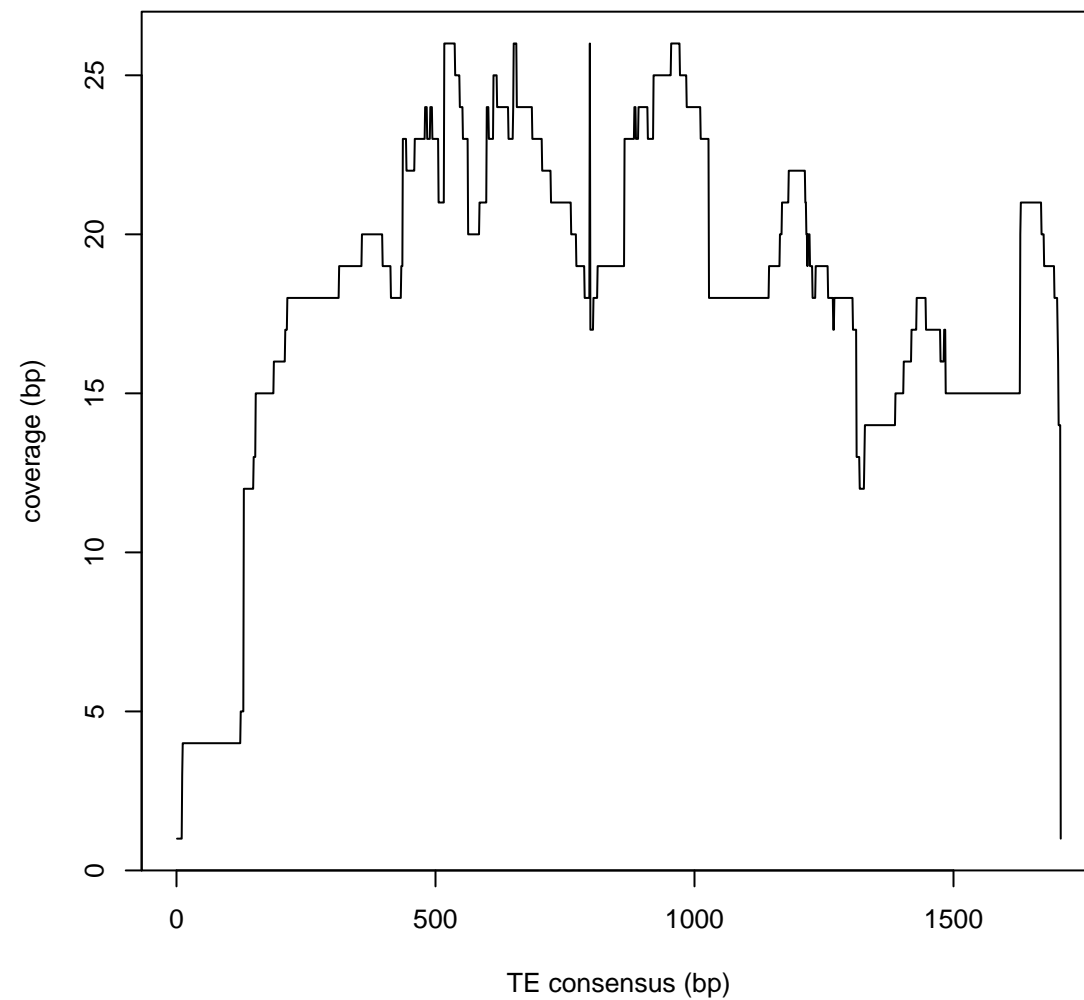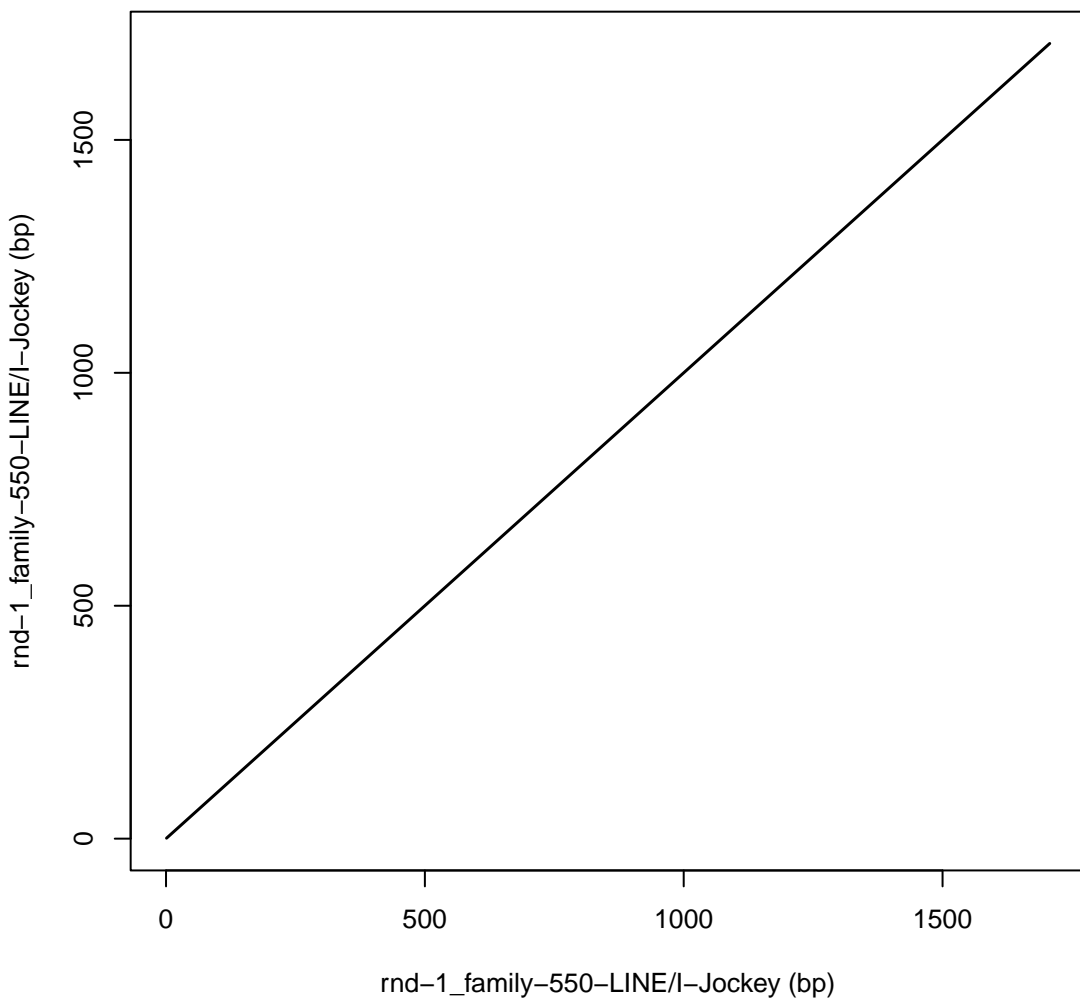**TE: rnd-1_family-550-LINE/I-Jockey**
consensus size: 1707bp; fragments: 82; full length: 1 (>=1536.3bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected