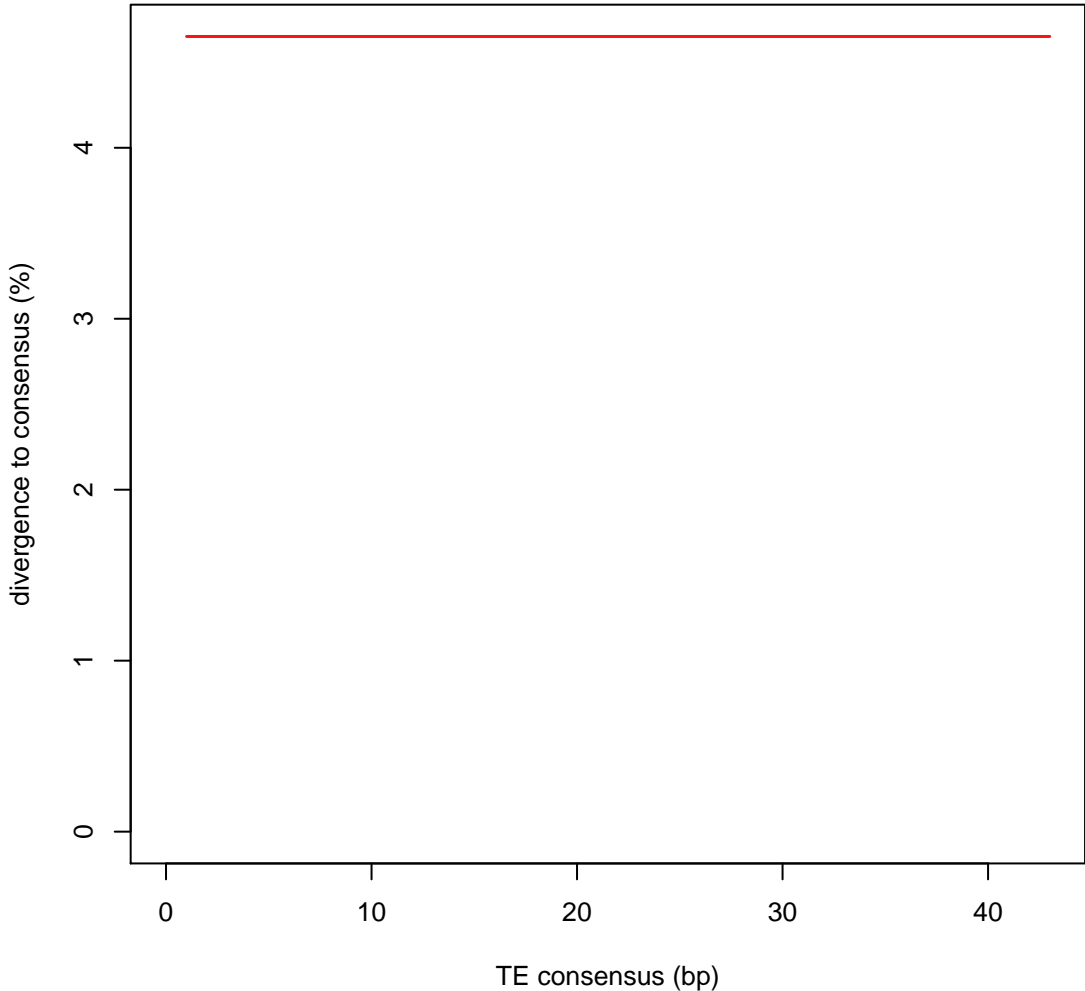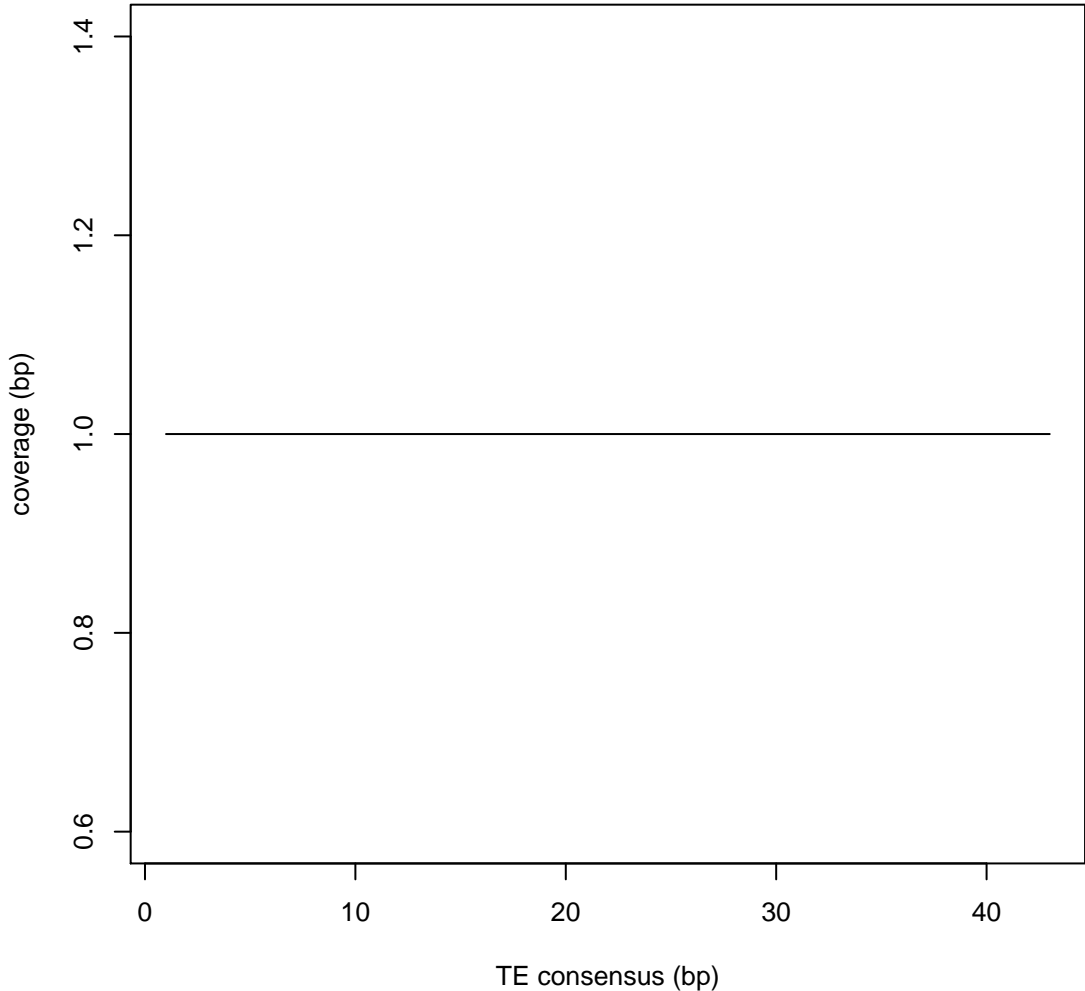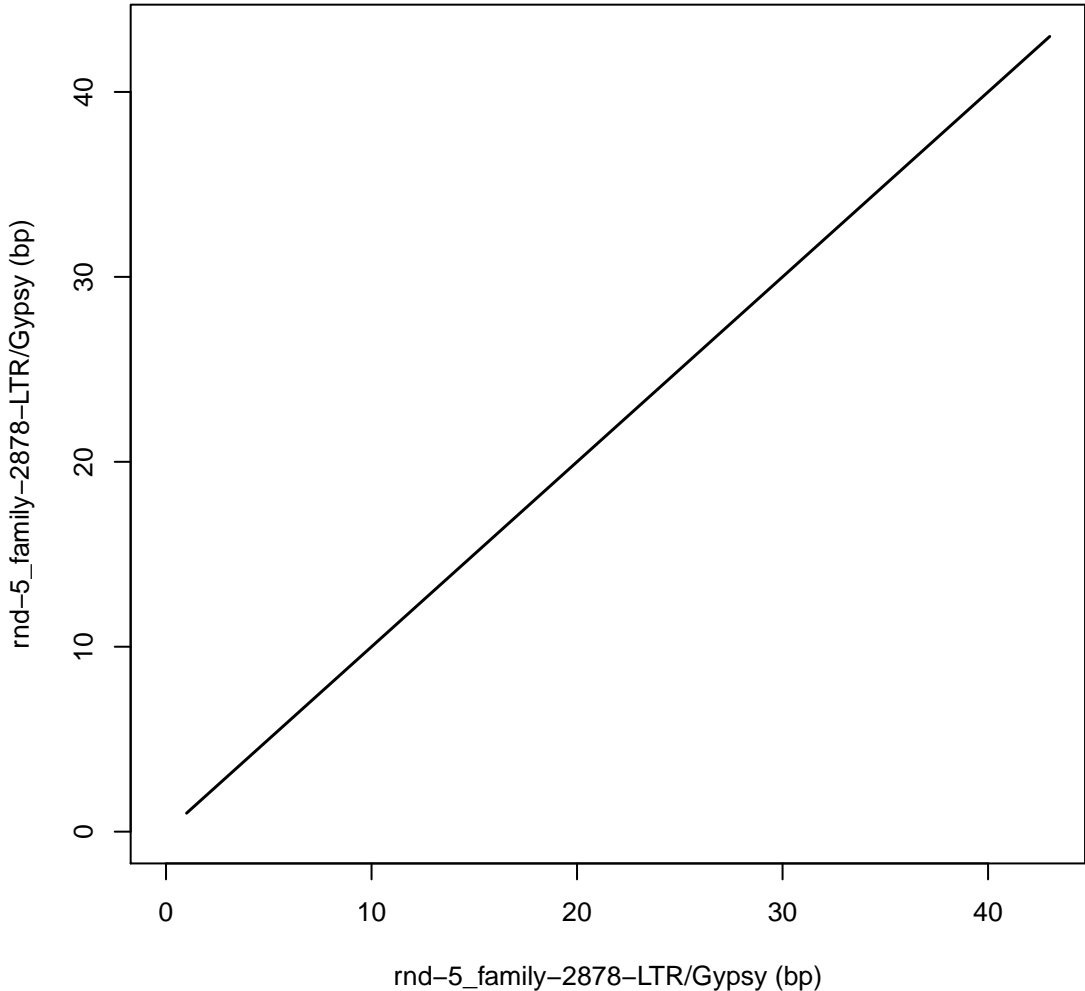**TE: rnd–5_family–2878–LTR/Gypsy**
consensus size: 43bp; fragments: 1; full length: 1 (>=38.7bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected