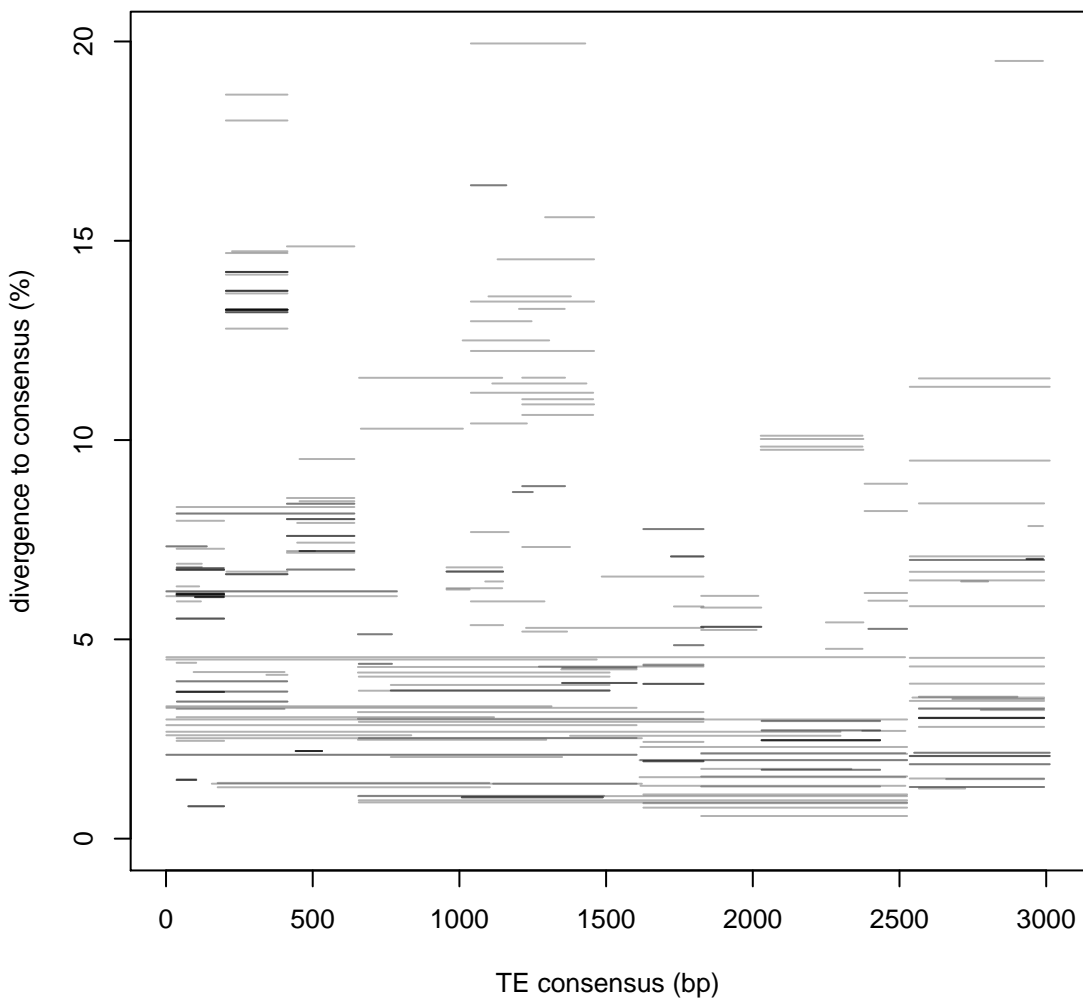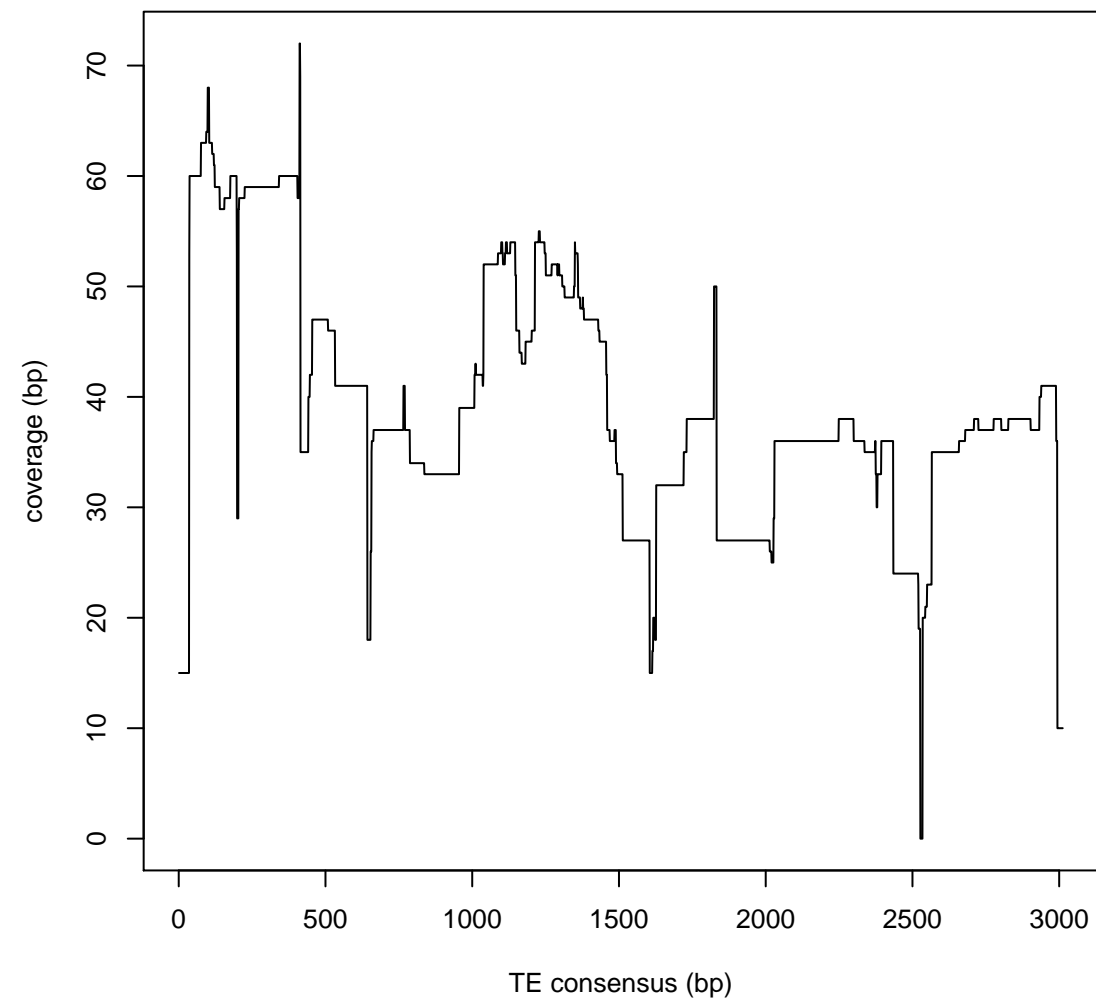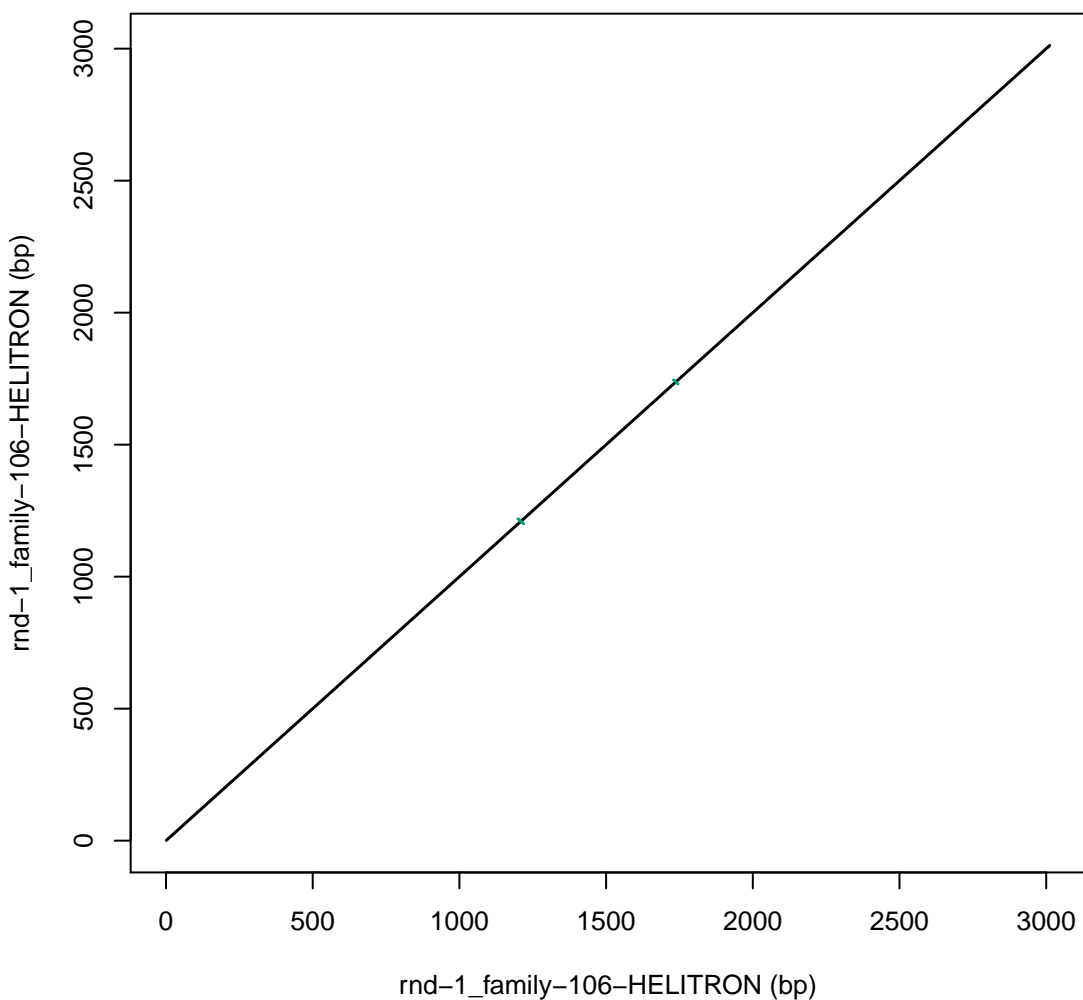**TE: rnd-1_family-106-HELITRON**
consensus size: 3012bp; fragments: 303; full length: 0 (>=2710.8bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected