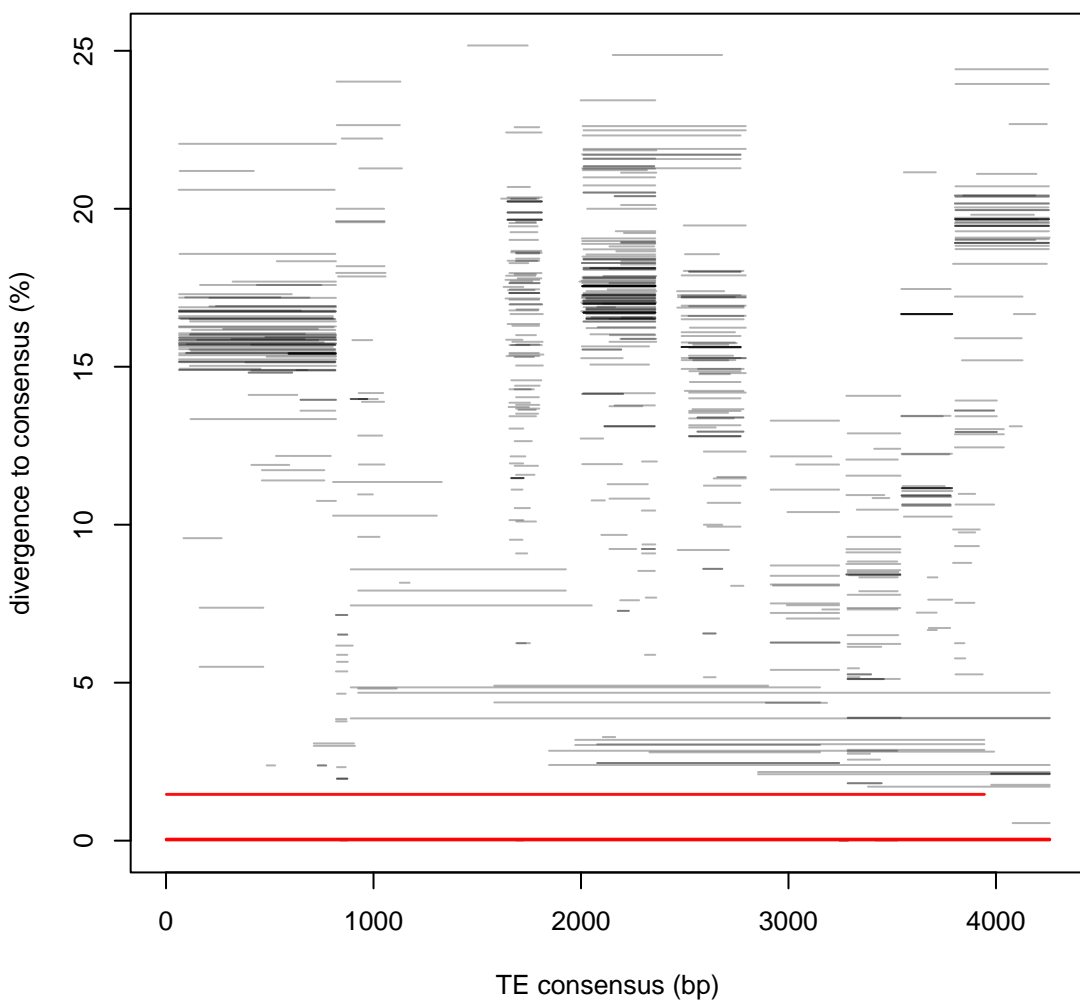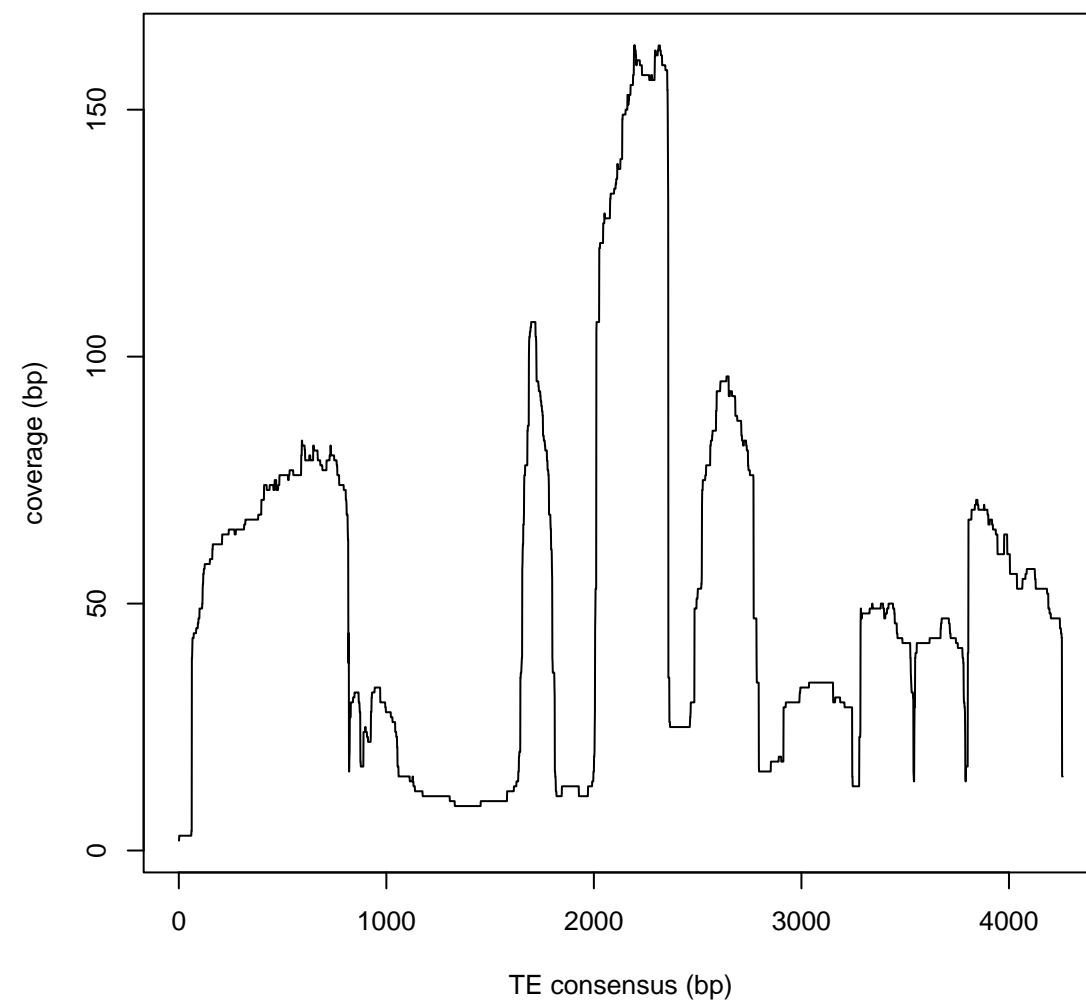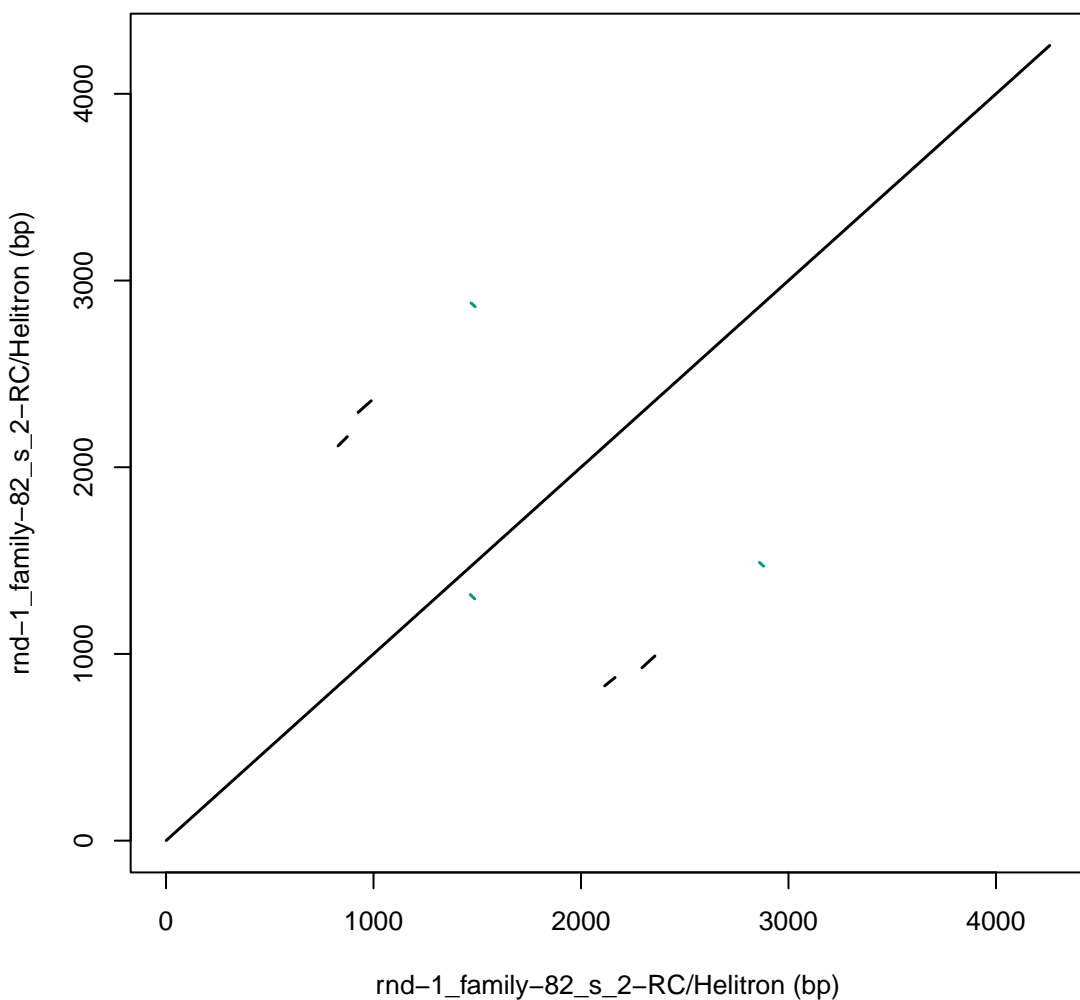**TE: rnd−1_family−82_s_2−RC/Helitron**
consensus size: 4259bp; fragments: 655; full length: 3 (>=3833.1bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected