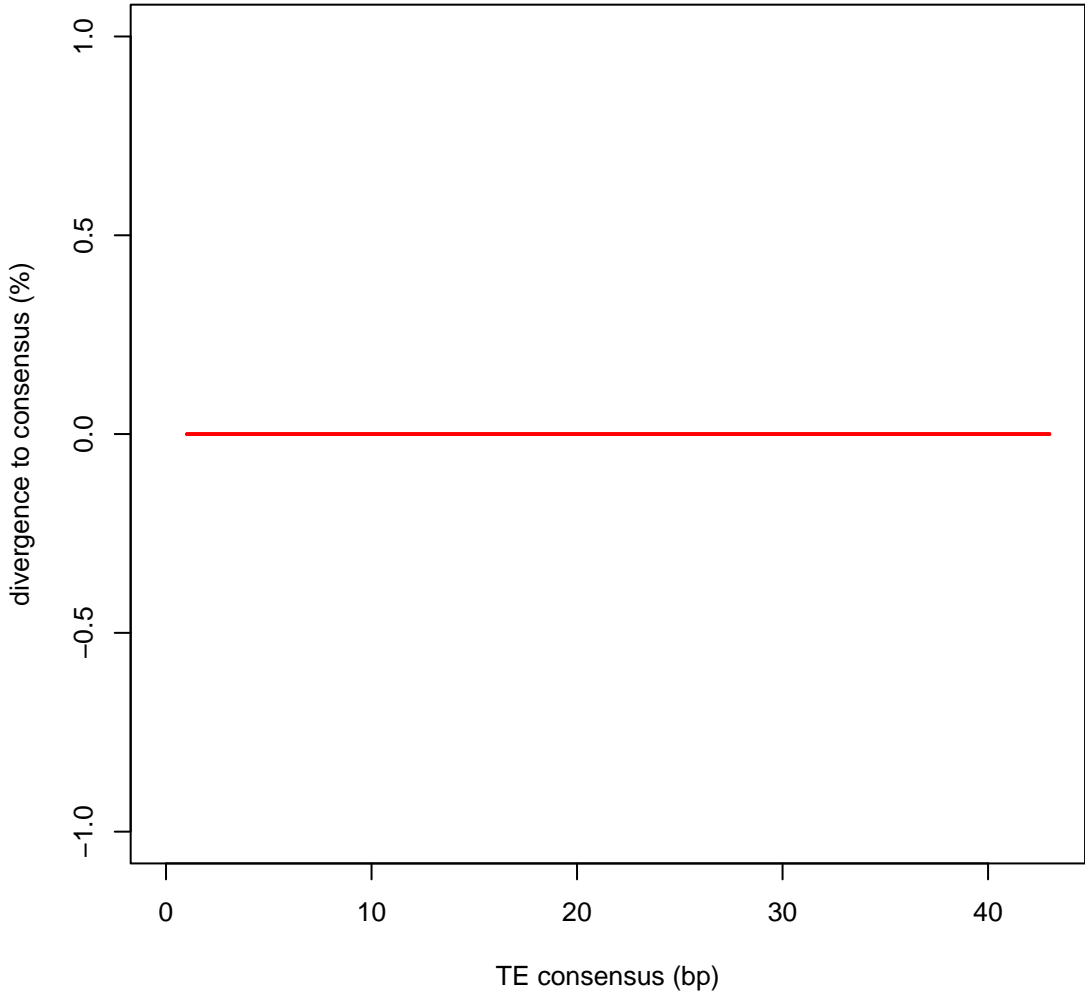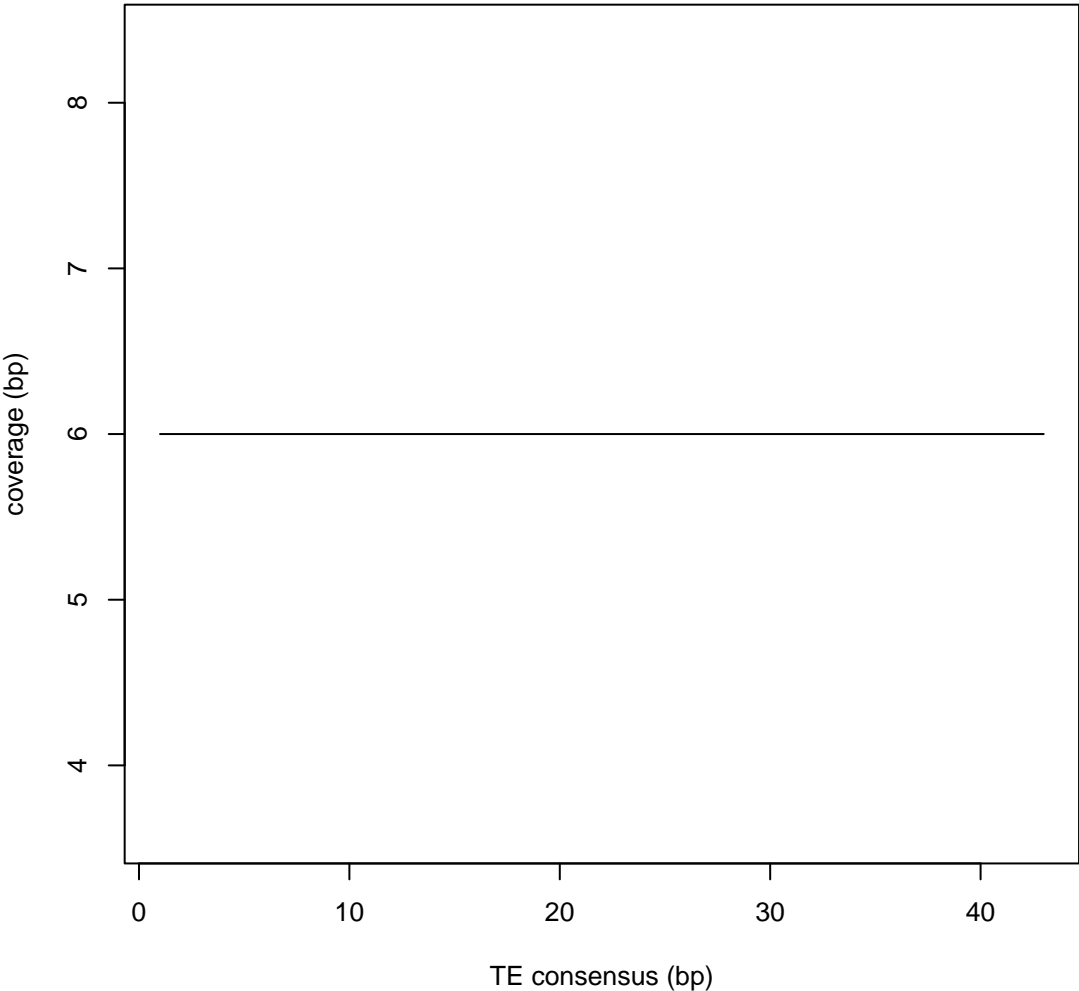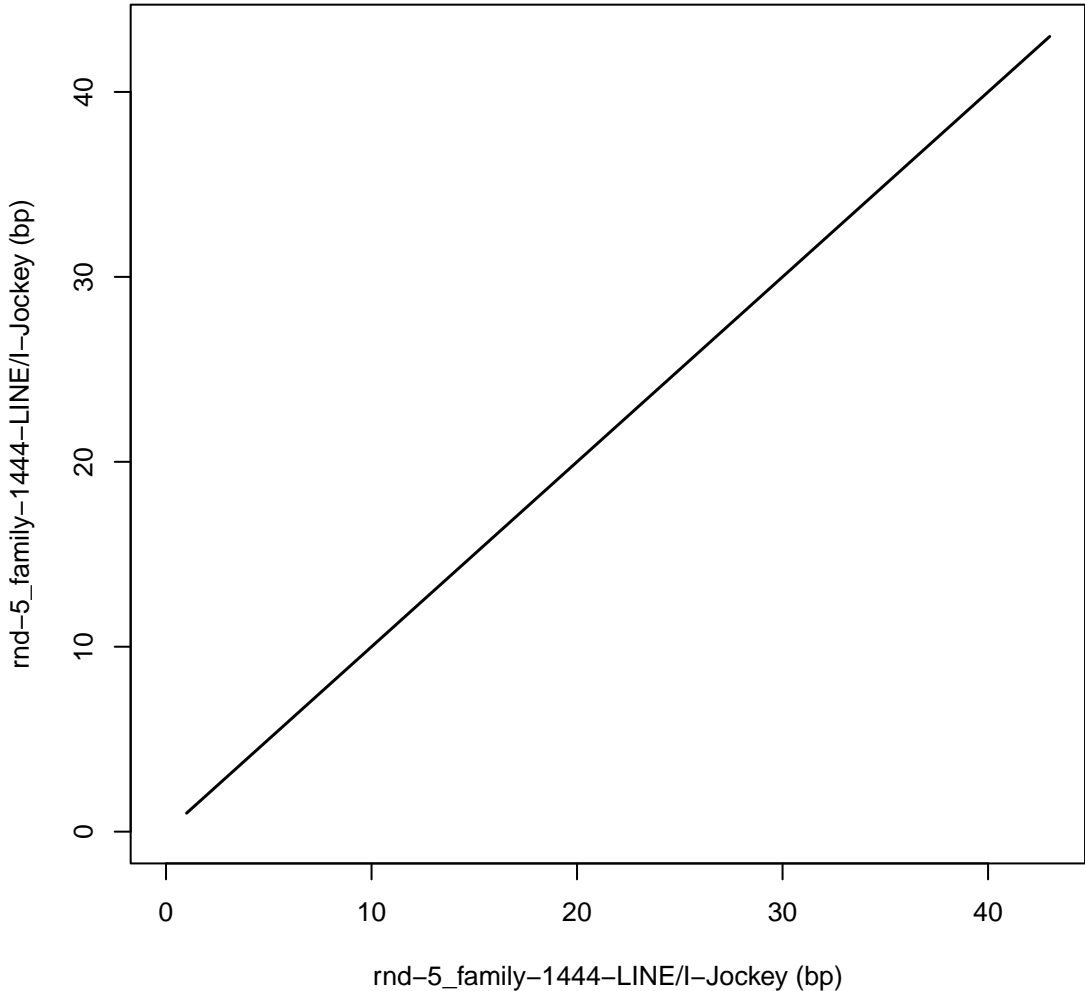**TE: rnd−5_family−1444−LINE/I−Jockey**
consensus size: 43bp; fragments: 6; full length: 6 (>=38.7bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected