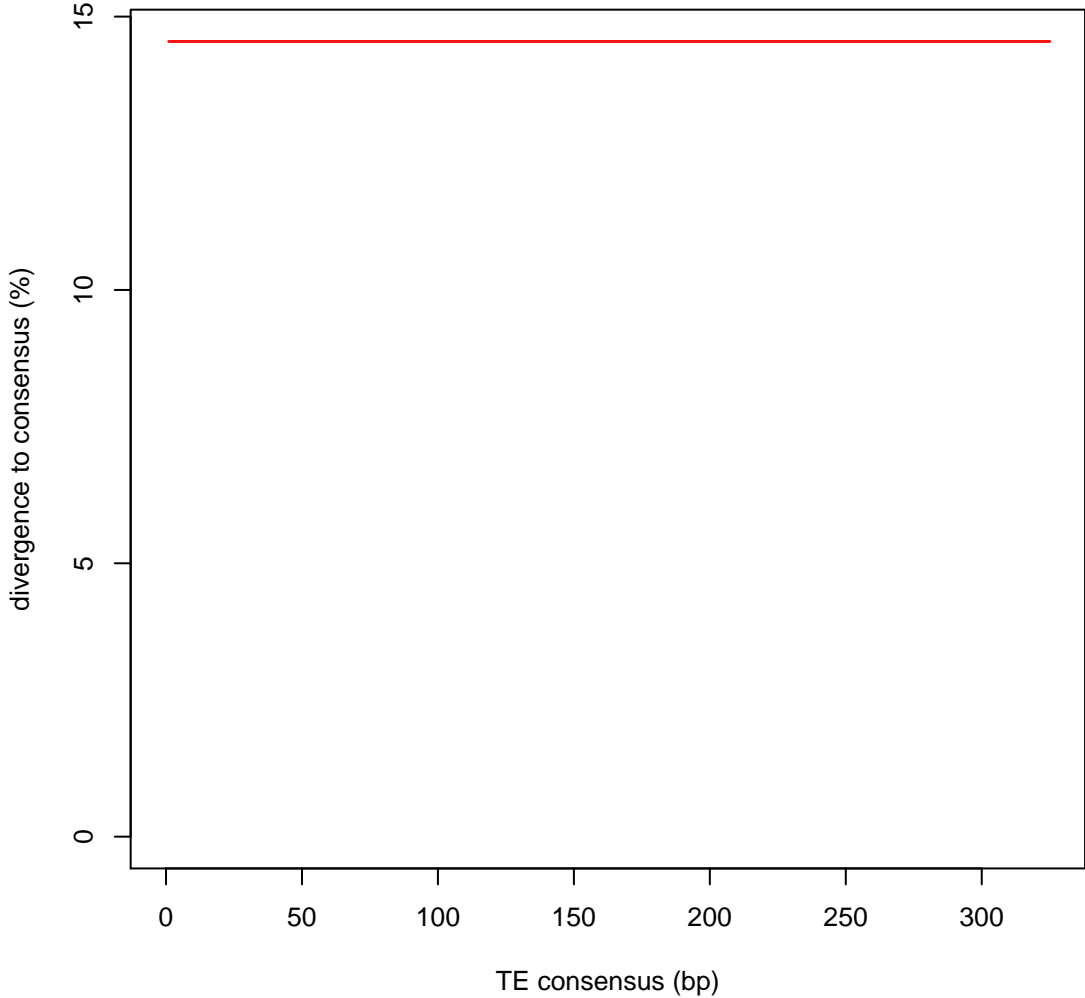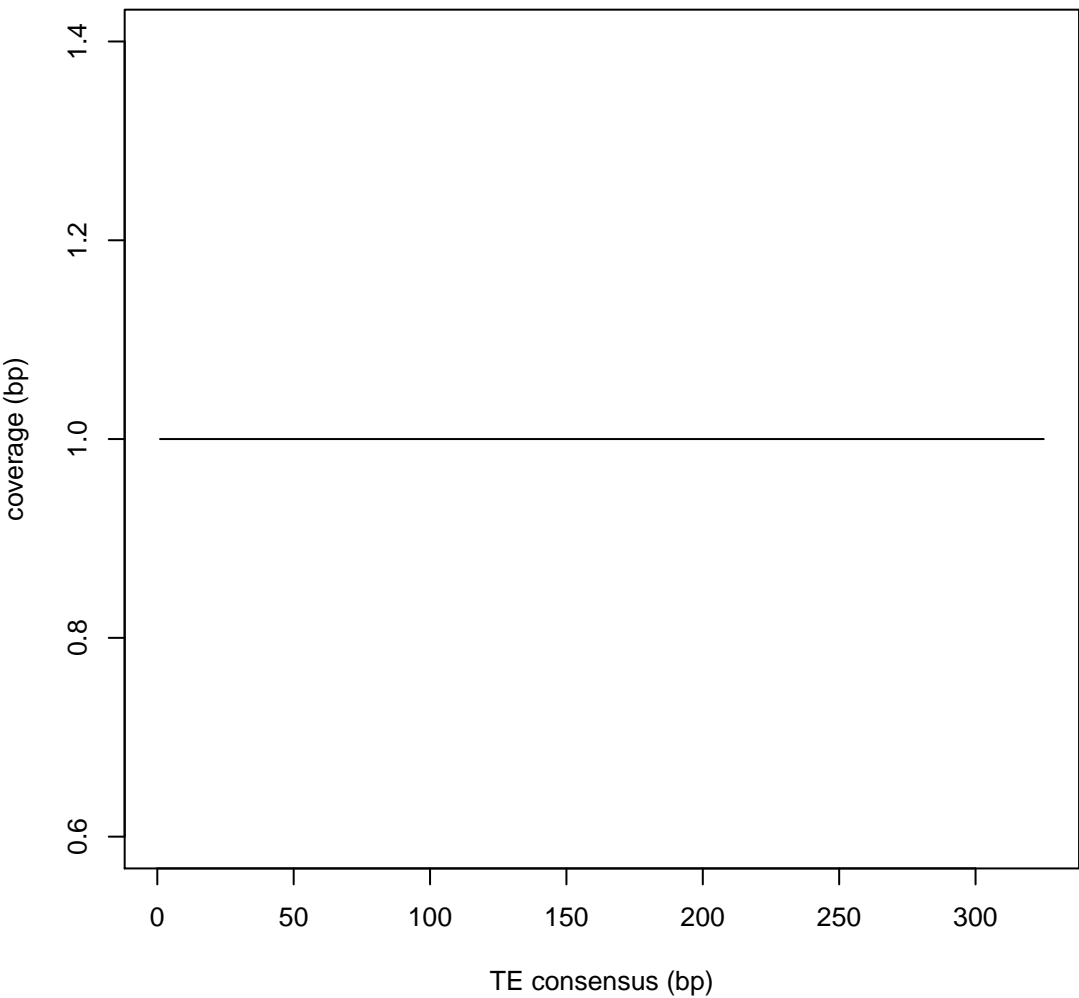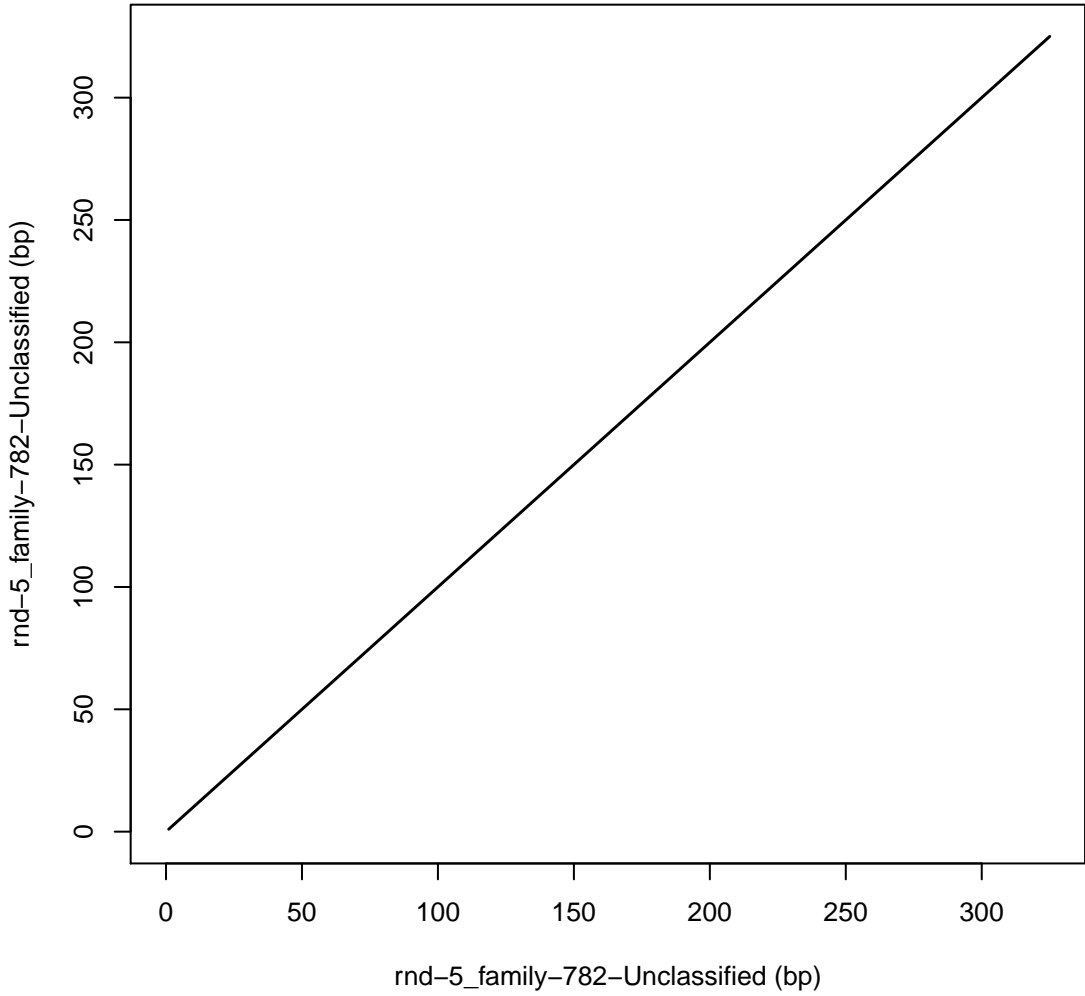**TE: rnd−5_family−782−Unclassified**
consensus size: 325bp; fragments: 1; full length: 1 (>=292.5bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected