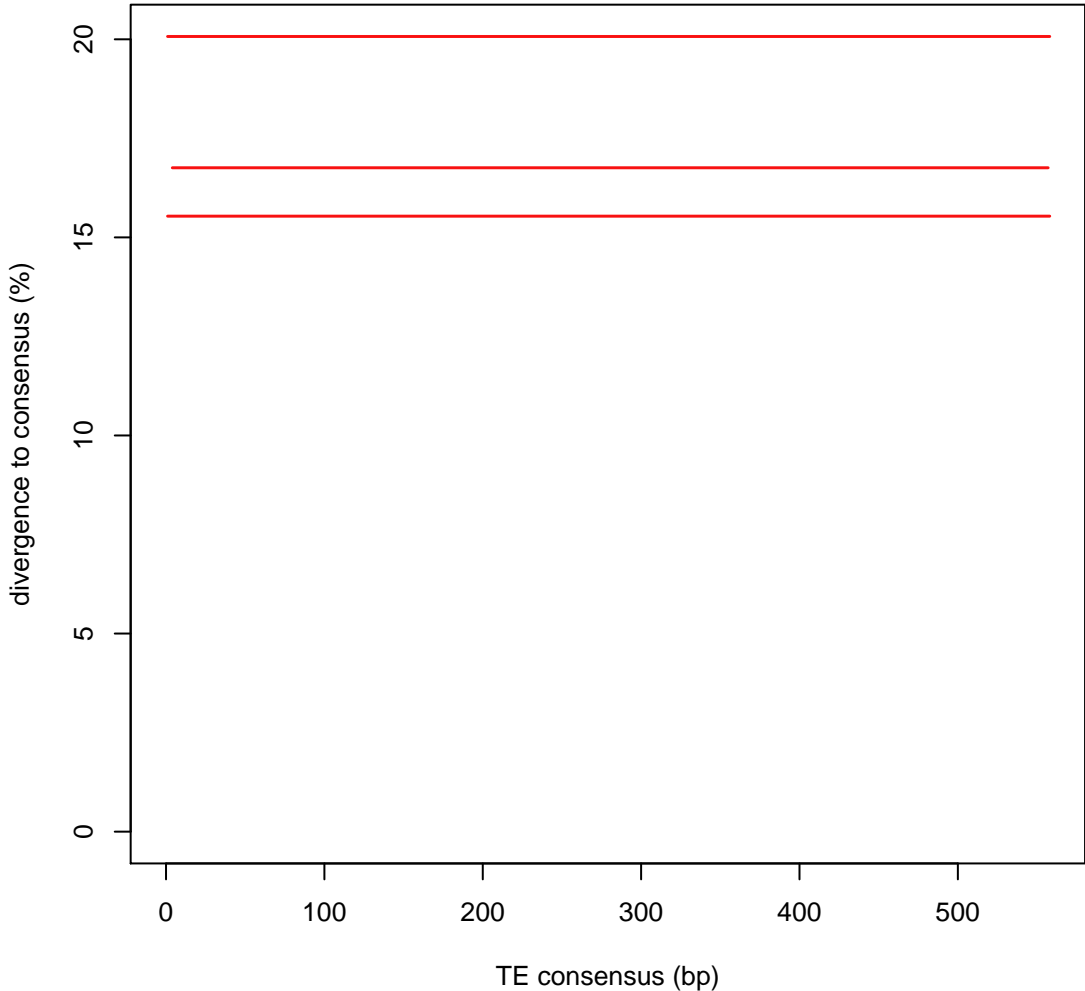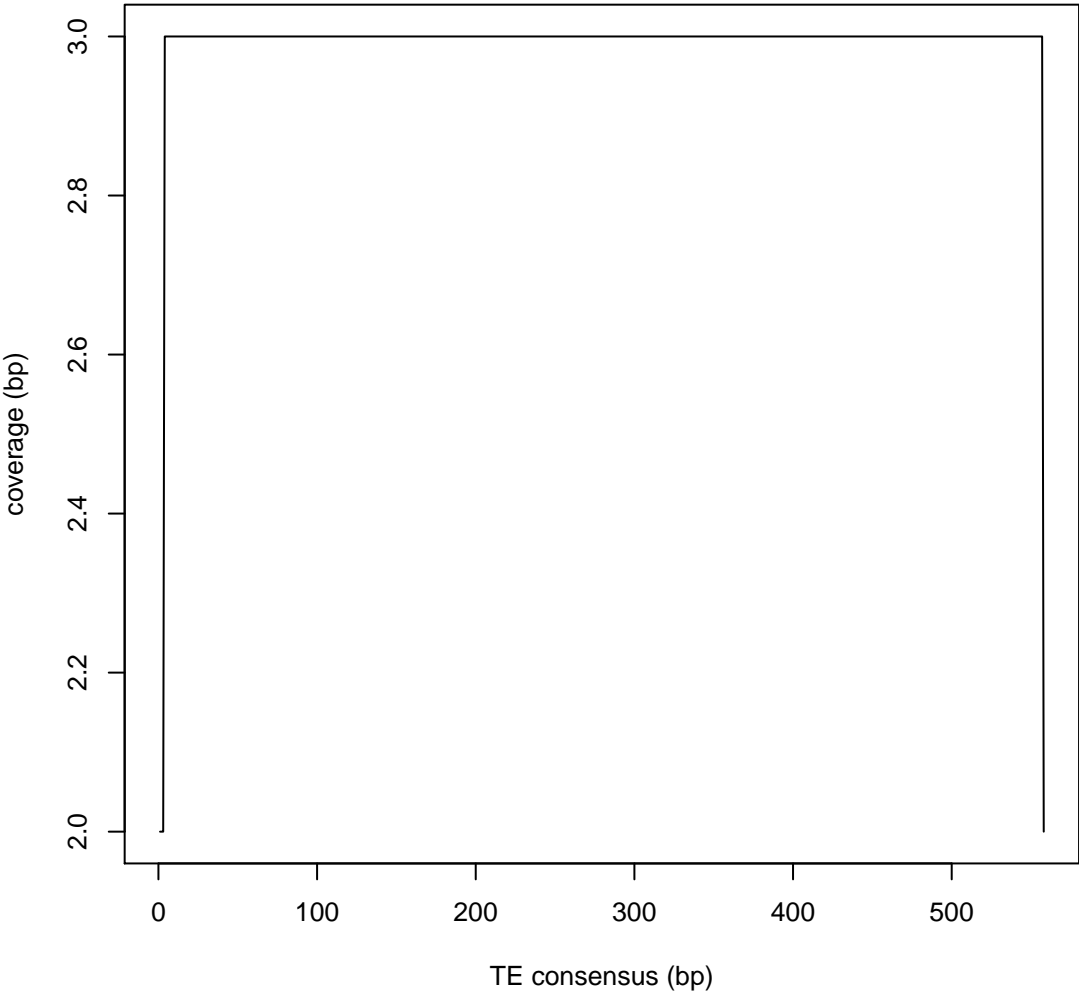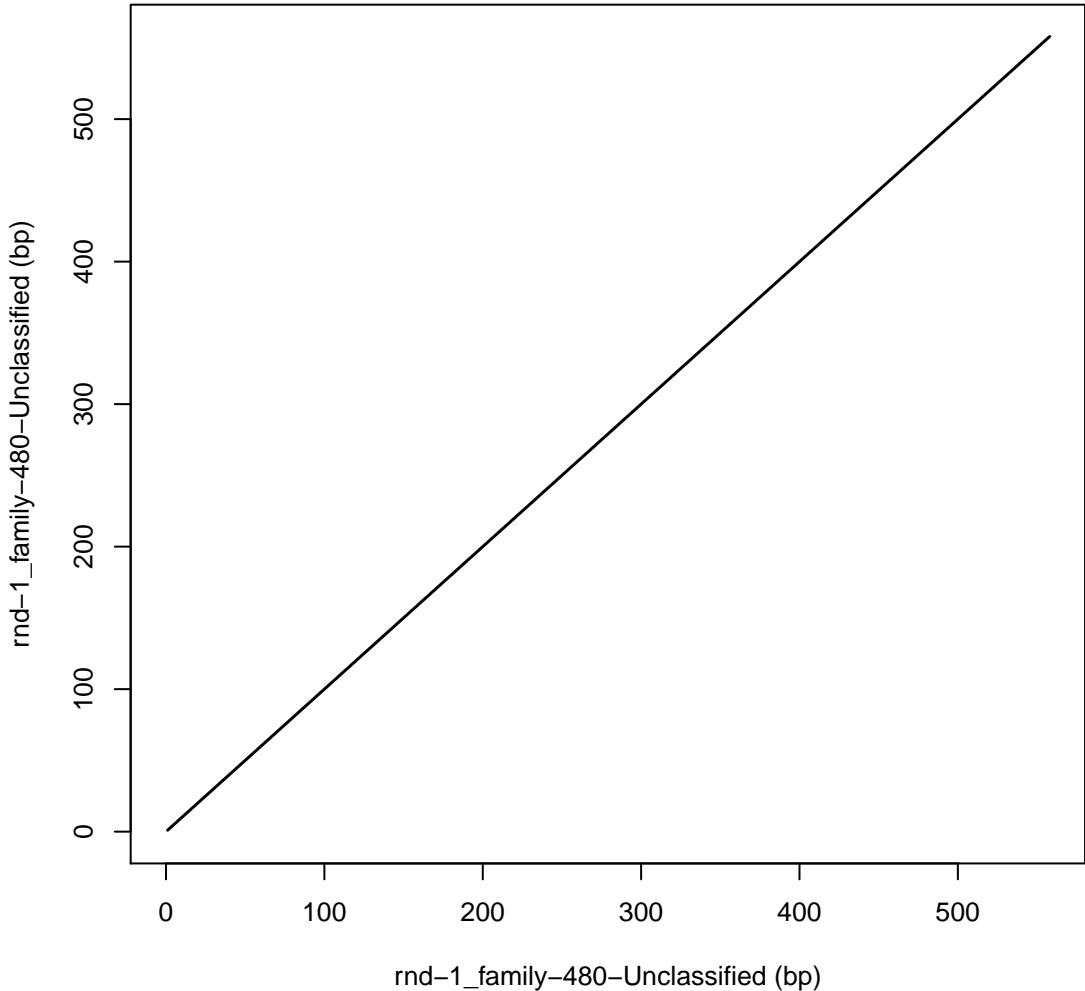**TE: rnd-1_family-480-Unclassified**
consensus size: 558bp; fragments: 3; full length: 3 (>=502.2bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected