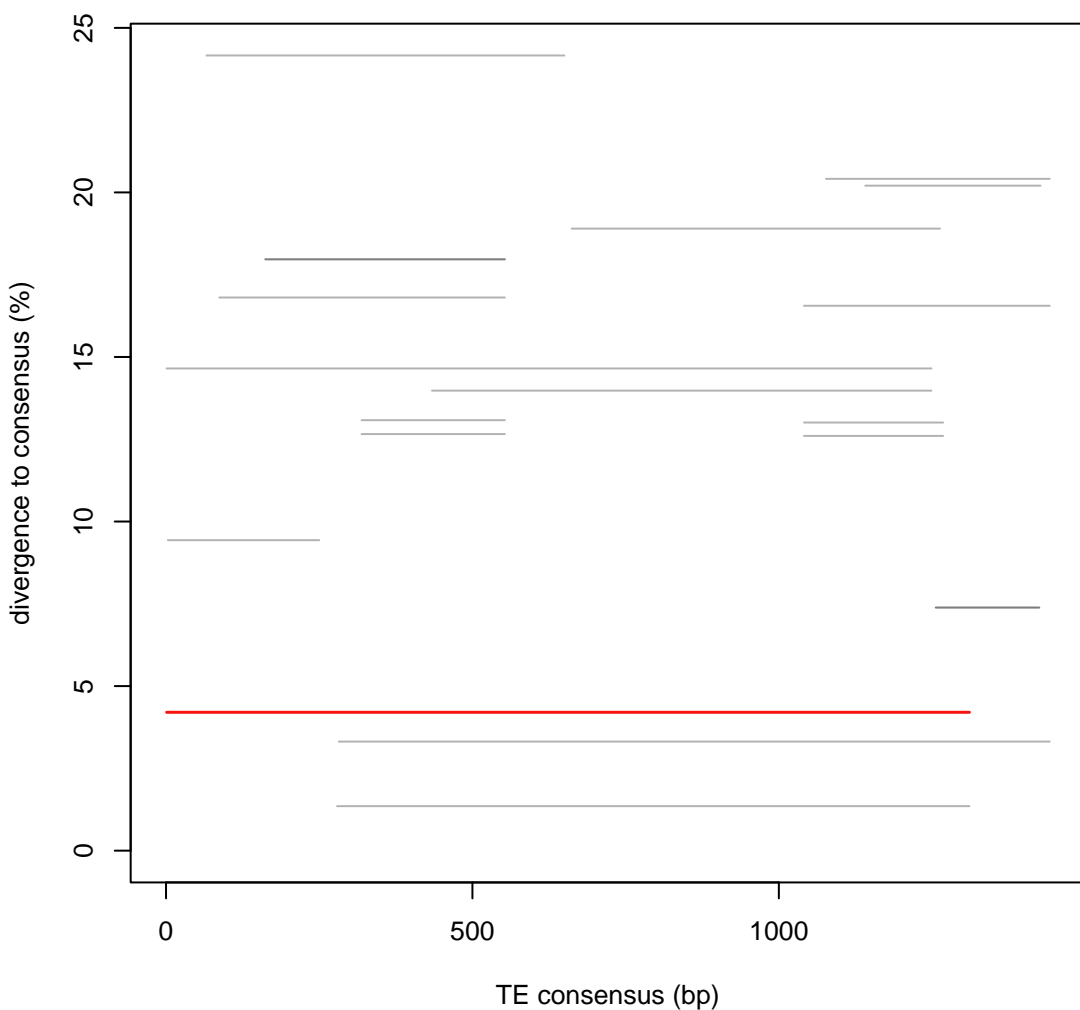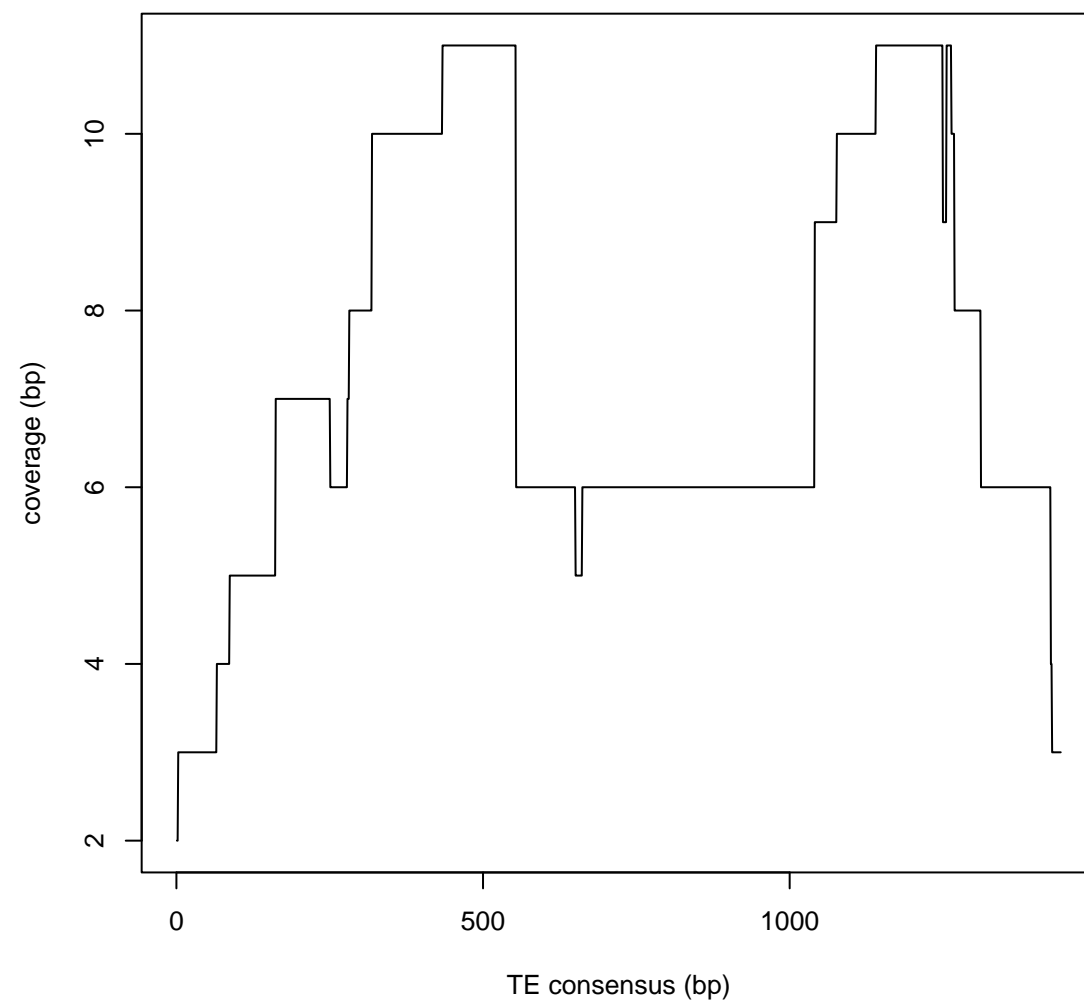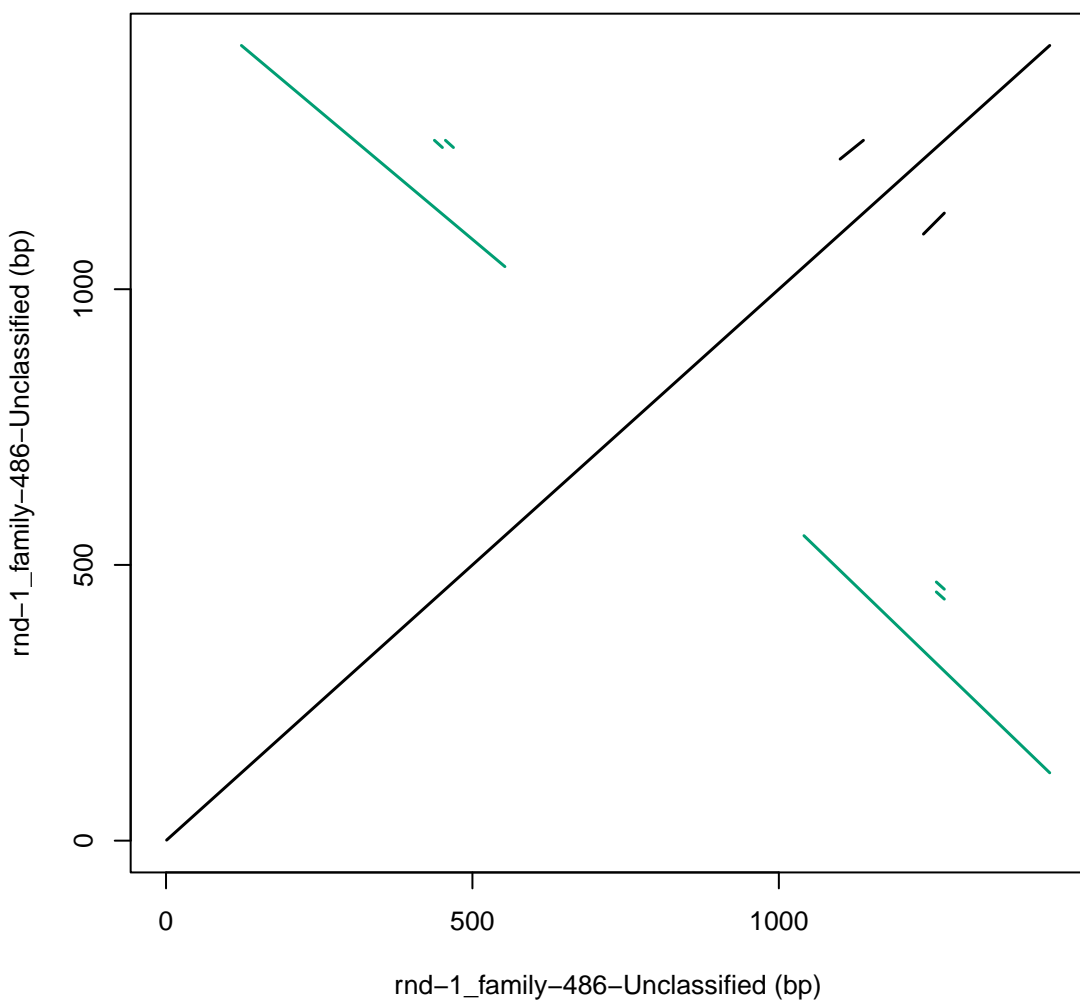**TE: rnd-1_family-486-Unclassified**
consensus size: 1442bp; fragments: 20; full length: 1 (>=1297.8bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected