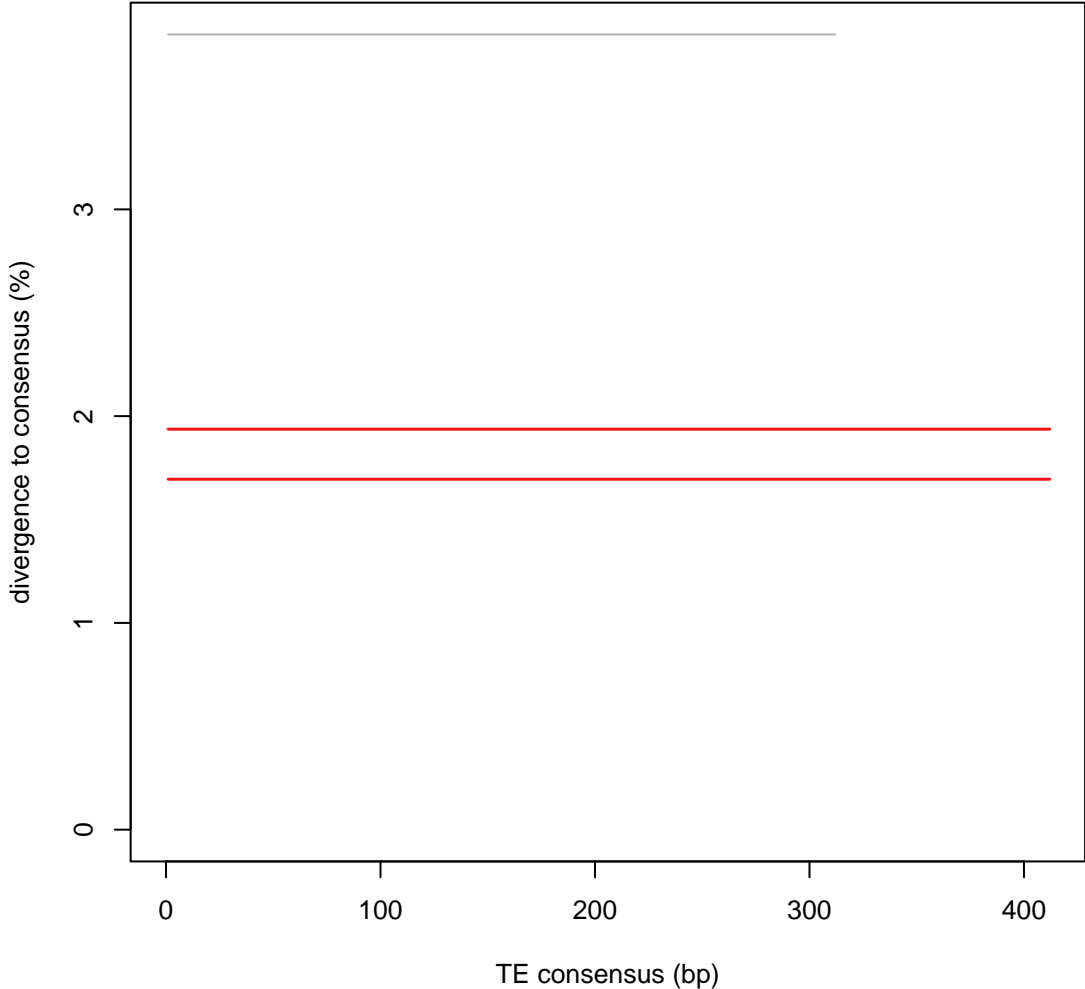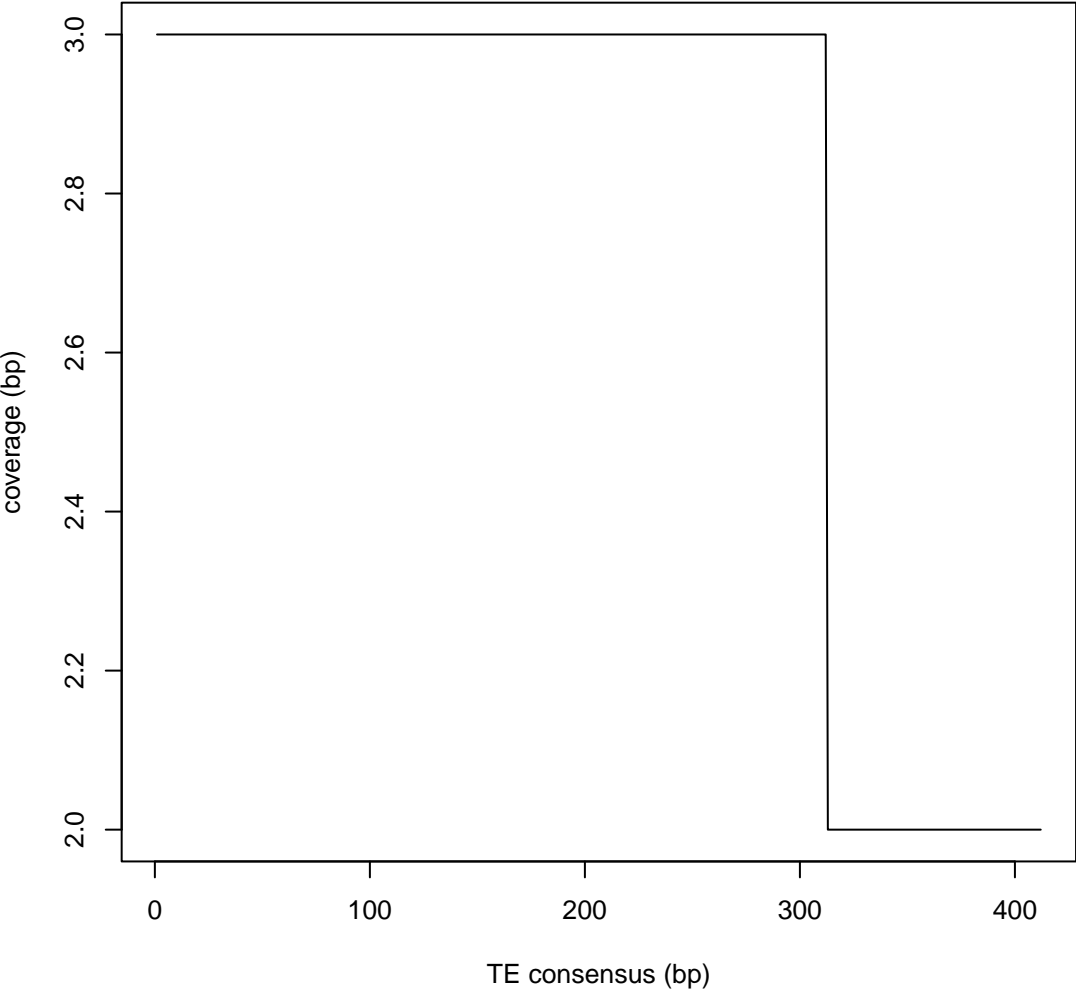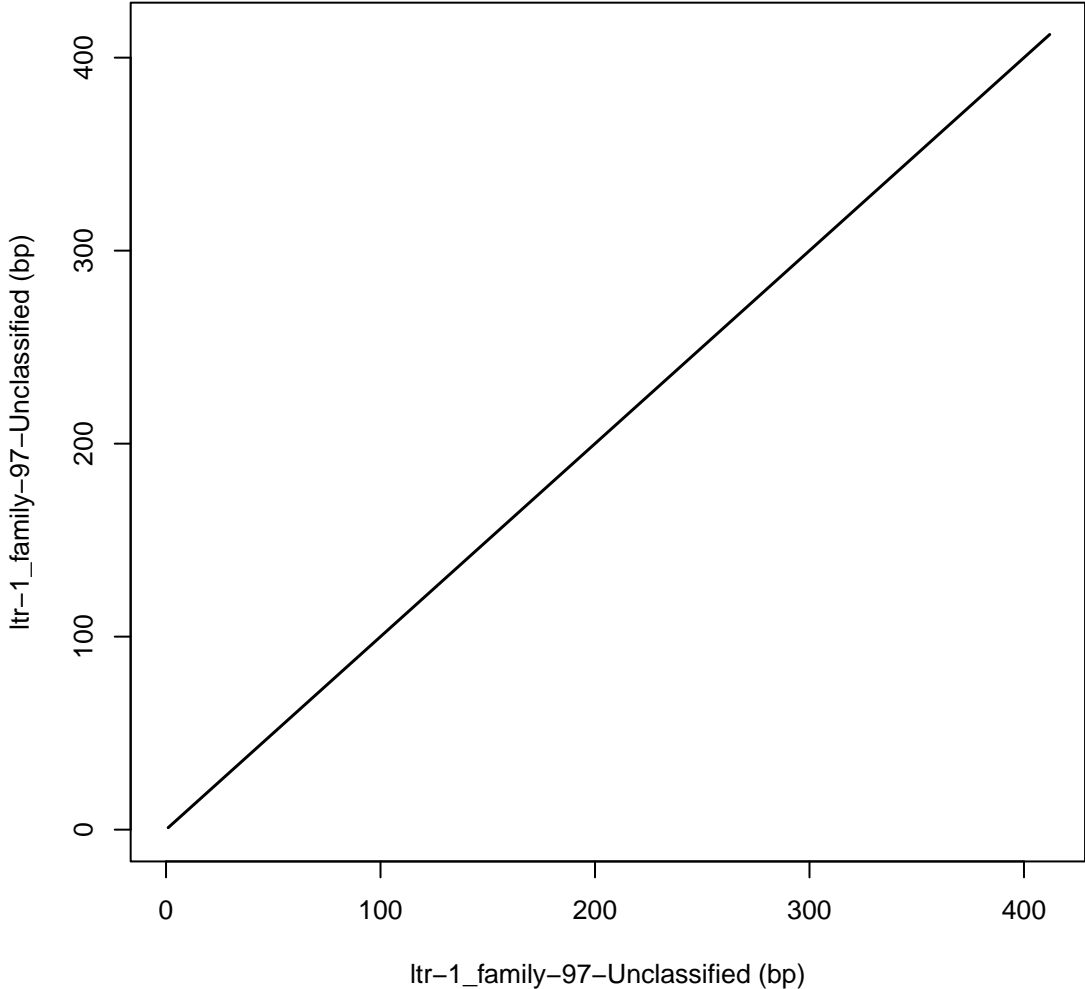**TE: ltr-1_family-97-Unclassified**
consensus size: 412bp; fragments: 3; full length: 2 (>=370.8bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected