TE: seq\_13167 consensus size: 2113bp; fragments: 2780; full length: 40 (>=1901.7bp) TE consensus genomic coverage divergence to consensus (%) coverage (bp) TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits seq\_13167 (bp) no orf >400 bp detected 

seq\_13167 (bp)

seq\_13167 (bp)