TE: seq_14442 TE consensus genomic coverage consensus size: 414bp; fragments: 1037; full length: 105 (>=372.6bp) divergence to consensus (%) coverage (bp) TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits seq_14442 (bp) no orf >400 bp detected seq_14442 (bp) seq_14442 (bp)