TE: seq\_21325 TE consensus genomic coverage consensus size: 2988bp; fragments: 2846; full length: 656 (>=2689.2bp) divergence to consensus (%) coverage (bp) TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits seq\_21325 (bp) 

seq\_21325 (bp)

no orf >400 bp detected

seq\_21325 (bp)