TE: seq_17220 TE consensus genomic coverage consensus size: 4220bp; fragments: 18203; full length: 2507 (>=3798bp) divergence to consensus (%) coverage (bp) TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits seq_17220 (bp) no orf >400 bp defected seq_17220 (bp) seq_17220 (bp)