TE: seq_21196 TE consensus genomic coverage consensus size: 4020bp; fragments: 21860; full length: 0 (>=3618bp) divergence to consensus (%) coverage (bp) TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits seq_21196 (bp) no orf >400 bp detected seq_21196 (bp) seq_21196 (bp)