TE: seq\_22677 TE consensus genomic coverage consensus size: 2427bp; fragments: 948; full length: 0 (>=2184.3bp) divergence to consensus (%) coverage (bp) TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits seq\_22677 (bp) no orf >400 bp detected seq\_22677 (bp) seq\_22677 (bp)