TE: seq\_17877 TE consensus genomic coverage consensus size: 2891bp; fragments: 54500; full length: 3086 (>=2601.9bp) divergence to consensus (%) coverage (bp) TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits JAM1\_pol LINE/RTE-BovB Copia-57\_BD LTR/Copia L2-2\_SP\_pol LINE/L2 seq\_17877 (bp) 

seq\_17877 (bp)

seq\_17877 (bp)