TE: rnd-1_family-479-TIR/TcMar-Pogo TE consensus genomic coverage consensus size: 2557bp; fragments: 122; full length: 0 (>=2301.3bp) divergence to consensus (%) coverage (bp) ∞ $^{\circ}$ TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits rnd-1_family-479-TIR/TcMar-Pogo (bp) UN-Pogo_tp TIR/TcMar-Pogo

rnd-1_family-479-TIR/TcMar-Pogo (bp)

rnd-1_family-479-TIR/TcMar-Pogo (bp)