TE: rnd-5_family-4576-TIR/TcMar-Tc1 TE consensus genomic coverage consensus size: 2495bp; fragments: 79; full length: 0 (>=2245.5bp) divergence to consensus (%) coverage (bp) TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits rnd-5_family-4576-TIR/TcMar-Tc1 (bp) Mariner2_DM_tp TIR/TcMar-Tc1

rnd-5_family-4576-TIR/TcMar-Tc1 (bp)

rnd-5_family-4576-TIR/TcMar-Tc1 (bp)