TE: rnd-1_family-513-LINE/I-Jockey TE consensus genomic coverage consensus size: 2472bp; fragments: 72; full length: 0 (>=2224.8bp) divergence to consensus (%) coverage (bp) TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits L1SU_SP_pol LINE/L1-Tx1 rnd-1_family-513-LINE/I-Jockey (bp) Gypsy-13_BG_pol LTR/Gyps G4_DM_pol LINE/I-Jockey G4_DM_pol LINE/I-Jockey

rnd-1_family-513-LINE/I-Jockey (bp)

rnd-1_family-513-LINE/I-Jockey (bp)