TE: rnd-1_family-161-LINE/I-Jockey TE consensus genomic coverage consensus size: 8693bp; fragments: 1741; full length: 4 (>=7823.7bp) divergence to consensus (%) coverage (bp) TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits rnd-1_family-161-LINE/I-Jockey (bp) key-6_DAn_pol LINE/I-Jockey Gypsy-9_PR_3p LTR/Gypsy Jockey-1_DGri_pol LINE/I-Jockey Gypsy-9_PR_3p LTI rnd-1_family-161-LINE/I-Jockey (bp) rnd-1_family-161-LINE/I-Jockey (bp)