TE: rnd-1_family-63-LINE consensus size: 4210bp; fragments: 1810; full length: 0 (>=3789bp) TE consensus genomic coverage 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-63-LINE (bp) Saci-2 LTR/Gypsy ^ℂGypsy−1_SCH_gag LTR/Gyp rnd-1_family-63-LINE (bp) rnd-1_family-63-LINE (bp)