TE: rnd-1\_family-63-LINE/I-Jockey TE consensus genomic coverage consensus size: 4140bp; fragments: 1825; full length: 1 (>=3726bp) 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/I-Jockey (bp) Saci-2 LTR/Gypsy Gypsy-1\_SCH\_gag LTR/Gyps rnd-1\_family-63-LINE/I-Jockey (bp) rnd-1\_family-63-LINE/I-Jockey (bp)