TE: rnd-1_family-219-LINE/I-Jockey TE consensus genomic coverage consensus size: 11214bp; fragments: 1887; full length: 3 (>=10092.6bp) 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-219-LINE/I-Jockey (bp) sy-9_PR_3p LTR/Gypsy □□ Jockey-1_DGri_pol LINE/I-Jockey Gypsy-9_PR_3p LTR/Gypsy Jockey-1_DGri_pol LINE/I-Jockey rnd-1_family-219-LINE/I-Jockey (bp) rnd-1_family-219-LINE/I-Jockey (bp)