TE: rnd-1_family-452_s_1-LTR/Gypsy consensus size: 11354bp; fragments: 1893; full length: 3 (>=10218.6bp) 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-452_s_1-LTR/Gypsy (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-452_s_1-LTR/Gypsy (bp)

rnd-1_family-452_s_1-LTR/Gypsy (bp)