TE: rnd-1_family-372-LINE TE consensus genomic coverage consensus size: 2634bp; fragments: 178; full length: 1 (>=2370.6bp) divergence to consensus (%) coverage (bp) TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits Jockey-7_DEu_pol LINE/I-Jockey rnd-1_family-372-LINE (bp) Gypsy-24_NV LTR/Gypsy

rnd-1_family-372-LINE (bp)

rnd-1_family-372-LINE (bp)