TE: rnd-1_family-372-LINE/I-Jockey TE consensus genomic coverage consensus size: 2629bp; fragments: 178; full length: 1 (>=2366.1bp) 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-372-LINE/I-Jockey (bp) no orf >400 bp detected

rnd-1_family-372-LINE/I-Jockey (bp)

rnd-1_family-372-LINE/I-Jockey (bp)