TE: rnd-1_family-513-LINE/I-Jockey TE consensus genomic coverage consensus size: 1583bp; fragments: 84; full length: 1 (>=1424.7bp) divergence to consensus (%) coverage (bp) TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits G2_DM_pol LINE/I-Jockey rnd-1_family-513-LINE/I-Jockey (bp)

rnd-1_family-513-LINE/I-Jockey (bp)

rnd-1_family-513-LINE/I-Jockey (bp)