TE: rnd-5\_family-3214-Unclassified TE consensus genomic coverage consensus size: 2185bp; fragments: 175; full length: 2 (>=1966.5bp) divergence to consensus (%) coverage (bp) TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits rnd-5\_family-3214-Unclassified (bp) no orf >400 bp detected 

rnd-5\_family-3214-Unclassified (bp)

rnd-5\_family-3214-Unclassified (bp)