TE: rnd-5_family-559-RC/Helitron TE consensus genomic coverage consensus size: 1027bp; fragments: 221; full length: 1 (>=924.3bp) 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-559-RC/Helitron (bp) no orf >400 bp detected

rnd-5_family-559-RC/Helitron (bp)

rnd-5_family-559-RC/Helitron (bp)