TE: rnd-5_family-3214-Unclassified TE consensus genomic coverage consensus size: 4613bp; fragments: 395; full length: 2 (>=4151.7bp) 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) Gypsy2_pol LTR/Gypsy BEL-599_AA_pol LTI rnd-5_family-3214-Unclassified (bp) rnd-5_family-3214-Unclassified (bp)