TE: rnd-1_family-350-LINE/CR1 TE consensus genomic coverage consensus size: 3049bp; fragments: 809; full length: 3 (>=2744.1bp) divergence to consensus (%) coverage (bp) TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits CR1-1_DRh_gag LINE/CR1 CR1-4_DEI_pol LINE/CR1 rnd-1_family-350-LINE/CR1 (bp) CR1-2_AG_pol LINE/CR1 rnd-1_family-350-LINE/CR1 (bp) rnd-1_family-350-LINE/CR1 (bp)