TE: rnd-1_family-252_s_2-LTR/Gypsy consensus size: 7480bp; fragments: 396; full length: 3 (>=6732bp) TE consensus genomic coverage divergence to consensus (%) coverage (bp) TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits rnd-1_family-252_s_2-LTR/Gypsy (bp) Gypsy-15_DEI_gag LTR/Gypsy Rover_DM_pol LTR/Gypsy Rover_DM_env LTR/Gypsy _ rnd-1_family-252_s_2-LTR/Gypsy (bp) rnd-1_family-252_s_2-LTR/Gypsy (bp)