TE: rnd-1_family-423-LTR/Gypsy consensus size: 2009bp; fragments: 54; full length: 0 (>=1808.1bp) 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-423-LTR/Gypsy (bp) Gypsy-24_DBp_pol LTR/Gypsy Gypsy-11_DTa_pol LTR/Gypsy

rnd-1_family-423-LTR/Gypsy (bp)

rnd-1_family-423-LTR/Gypsy (bp)