TE: rnd-5_family-370-LTR/Gypsy consensus size: 1863bp; fragments: 111; full length: 7 (>=1676.7bp) 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 ypsy37_AG LTR/Gypsy BEL-4_NV_pol LTR/Pao rnd-5_family-370-LTR/Gypsy (bp)

rnd-5_family-370-LTR/Gypsy (bp)

rnd-5_family-370-LTR/Gypsy (bp)