TE: ltr-1_family-63_s_1_s_1_s_2-LTR/Gypsy TE consensus genomic coverage consensus size: 7213bp; fragments: 308; full length: 3 (>=6491.7bp) divergence to consensus (%) coverage (bp) TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits ltr-1_family-63_s_1_s_1_s_2-LTR/Gypsy (bp) Gypsy-15_DEI_gag LTR/Gypsy Rover_DM_pol LTR/Gypsy Rover_DM_env LTR/Gypsy

ltr-1_family-63_s_1_s_1_s_2-LTR/Gypsy (bp)

ltr-1_family-63_s_1_s_1_s_2-LTR/Gypsy (bp)