TE: ltr-1\_family-63\_s\_1\_s\_1\_s\_1-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\_1-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\_1-LTR/Gypsy (bp)

ltr-1\_family-63\_s\_1\_s\_1\_s\_1-LTR/Gypsy (bp)