TE: Dmel-B-G971-Map3-LINE/nan TE consensus genomic coverage consensus size: 4123bp; fragments: 749; full length: 0 (>=3710.7bp) divergence to consensus (%) coverage (bp) TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits Dmel-B-G971-Map3-LINE/nan (bp) y5_pol LTR/Gypsy DMCR1A_gag LINE/CR1

Dmel-B-G971-Map3-LINE/nan (bp)

Dmel-B-G971-Map3-LINE/nan (bp)