TE: Dmel-B-R1653-Map3-LINE/nan TE consensus genomic coverage consensus size: 2505bp; fragments: 84; full length: 2 (>=2254.5bp) divergence to consensus (%) coverage (bp) TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits Dmel-B-R1653-Map3-LINE/nan (bp) no orf >400 bp detected 

Dmel-B-R1653-Map3-LINE/nan (bp)

Dmel-B-R1653-Map3-LINE/nan (bp)