TE: Dmel-B-R113-Map20-CLASSI/LINE/nan TE consensus genomic coverage consensus size: 2622bp; fragments: 381; full length: 1 (>=2359.8bp) divergence to consensus (%) coverage (bp) TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits × × Dmel-B-R113-Map20-CLASSI/LINE/nan (bp) R1-4_AP_pol LINE/R1 LOA_pol LINE/R1-LOA LOA-8_AAe_pol LINE/R1-LO

Dmel-B-R113-Map20-CLASSI/LINE/nan (bp)

Dmel-B-R113-Map20-CLASSI/LINE/nan (bp)