TE: Dmel-B-R301-Map9_reversed-LINE TE consensus genomic coverage consensus size: 4580bp; fragments: 119; full length: 7 (>=4122bp) divergence to consensus (%) coverage (bp) TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits > Dmel-B-R301-Map9_reversed-LINE (bp) HETA_gag LINE/I-Jockey Gypsy-97_MLP_pol LTR/Gypsy

Dmel-B-R301-Map9_reversed-LINE (bp)

Dmel-B-R301-Map9_reversed-LINE (bp)