TE: Dmel-B-R193-Map3-LINE/nan TE consensus genomic coverage consensus size: 2509bp; fragments: 166; full length: 2 (>=2258.1bp) divergence to consensus (%) coverage (bp) TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits DOC4_DM_gag LINE/I-Jockey Dmel-B-R193-Map3-LINE/nan (bp) Dmel-B-R193-Map3-LINE/nan (bp) Dmel-B-R193-Map3-LINE/nan (bp)