TE: Dmel-B-R193-Map3-LINE TE consensus genomic coverage consensus size: 2291bp; fragments: 166; full length: 2 (>=2061.9bp) divergence to consensus (%) coverage (bp) TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits FW2_DM_pol LINE/I-Jockey Dmel-B-R193-Map3-LINE (bp) Dmel-B-R193-Map3-LINE (bp) Dmel-B-R193-Map3-LINE (bp)