TE: Dmel-B-G1851-Map13-UNCLASSIFIED TE consensus genomic coverage consensus size: 4114bp; fragments: 385; full length: 7 (>=3702.6bp) divergence to consensus (%) coverage (bp) TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits × Dmel-B-G1851-Map13-UNCLASSIFIED (bp) Pifo\_env LTR/Gypsy 

Dmel-B-G1851-Map13-UNCLASSIFIED (bp)

Dmel-B-G1851-Map13-UNCLASSIFIED (bp)