TE: Dmel-B-P24-CLASSII/TIR/nan TE consensus genomic coverage consensus size: 1467bp; fragments: 120; full length: 4 (>=1320.3bp) divergence to consensus (%) coverage (bp) TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits Dmel-B-P24-CLASSII/TIR/nan (bp) Transib-4\_DBp TIR/CMC-Transib 

Dmel-B-P24-CLASSII/TIR/nan (bp)

Dmel-B-P24-CLASSII/TIR/nan (bp)