TE: Dmel-B-R4-Map3-LTR/Gypsy consensus size: 2195bp; fragments: 169; full length: 0 (>=1975.5bp) TE consensus genomic coverage divergence to consensus (%) coverage (bp) TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits Dmel-B-R4-Map3-LTR/Gypsy (bp) no orf >400 bp detected 

Dmel-B-R4-Map3-LTR/Gypsy (bp)

Dmel-B-R4-Map3-LTR/Gypsy (bp)