TE: Dmel-B-G2604-Map8-CLASSI/LTR/Gypsy TE consensus genomic coverage consensus size: 1182bp; fragments: 264; full length: 9 (>=1063.8bp) divergence to consensus (%) coverage (bp) TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits Dmel-B-G2604-Map8-CLASSI/LTR/Gypsy (bp) no orf >400 bp detected 

Dmel-B-G2604-Map8-CLASSI/LTR/Gypsy (bp)

Dmel-B-G2604-Map8-CLASSI/LTR/Gypsy (bp)