TE: Dmel-B-G1258-Map5_reversed-LTR/Gypsy TE consensus genomic coverage consensus size: 4310bp; fragments: 12241; full length: 0 (>=3879bp) divergence to consensus (%) coverage (bp) TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits Dmel-B-G1258-Map5_reversed-LTR/Gypsy (bp) no orf >400 bp detected Dmel-B-G1258-Map5_reversed-LTR/Gypsy (bp) Dmel-B-G1258-Map5_reversed-LTR/Gypsy (bp)