TE: Dmel-B-R4-Map3-LTR/GYPSY TE consensus genomic coverage consensus size: 2192bp; fragments: 169; full length: 0 (>=1972.8bp) divergence to consensus (%) coverage (bp) TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits Gypsy3_env LTR/Gypsy Dmel-B-R4-Map3-LTR/GYPSY (bp) Dmel-B-R4-Map3-LTR/GYPSY (bp) Dmel-B-R4-Map3-LTR/GYPSY (bp)