TE: Dmel-B-R1188-Map3-LTR/Gypsy consensus size: 4058bp; fragments: 219; full length: 4 (>=3652.2bp) 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 Gypsy5_pol LTR/Gypsy Dmel-B-R1188-Map3-LTR/Gypsy (bp) DMCR1A_gag LINE/CR1 Dmel-B-R1188-Map3-LTR/Gypsy (bp) Dmel-B-R1188-Map3-LTR/Gypsy (bp)