TE: Dmel-B-R1188-Map3-LTR/GYPSY TE consensus genomic coverage consensus size: 4082bp; fragments: 219; full length: 4 (>=3673.8bp) divergence to consensus (%) coverage (bp) TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits Gypsy5_Dpse_pol LTR/Gypsy Dmel-B-R1188-Map3-LTR/GYPSY (bp) ACCORD2_pol LTR/Gypsy CR1-1_DRh_gag LINE/CR1 Dmel-B-R1188-Map3-LTR/GYPSY (bp) Dmel-B-R1188-Map3-LTR/GYPSY (bp)