TE: Dmel-L-B236-Map1-CLASSI/LTR/Gypsy TE consensus genomic coverage consensus size: 5857bp; fragments: 593; full length: 1 (>=5271.3bp) divergence to consensus (%) coverage (bp) TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits Dmel-L-B236-Map1-CLASSI/LTR/Gypsy (bp) QUASIMODO_pol LTR/Gypsy QUASIMODO_env LTR/Gypsy

Dmel-L-B236-Map1-CLASSI/LTR/Gypsy (bp)

Dmel-L-B236-Map1-CLASSI/LTR/Gypsy (bp)