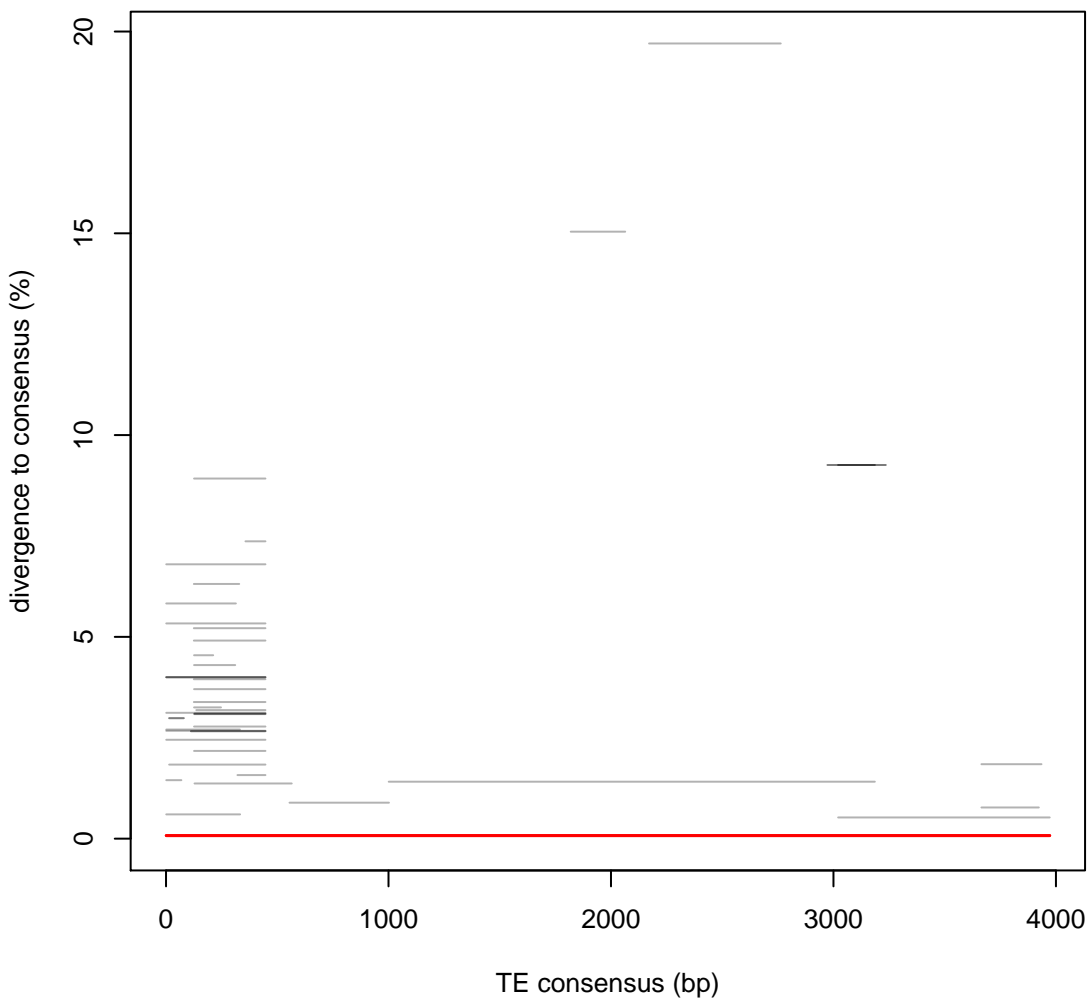
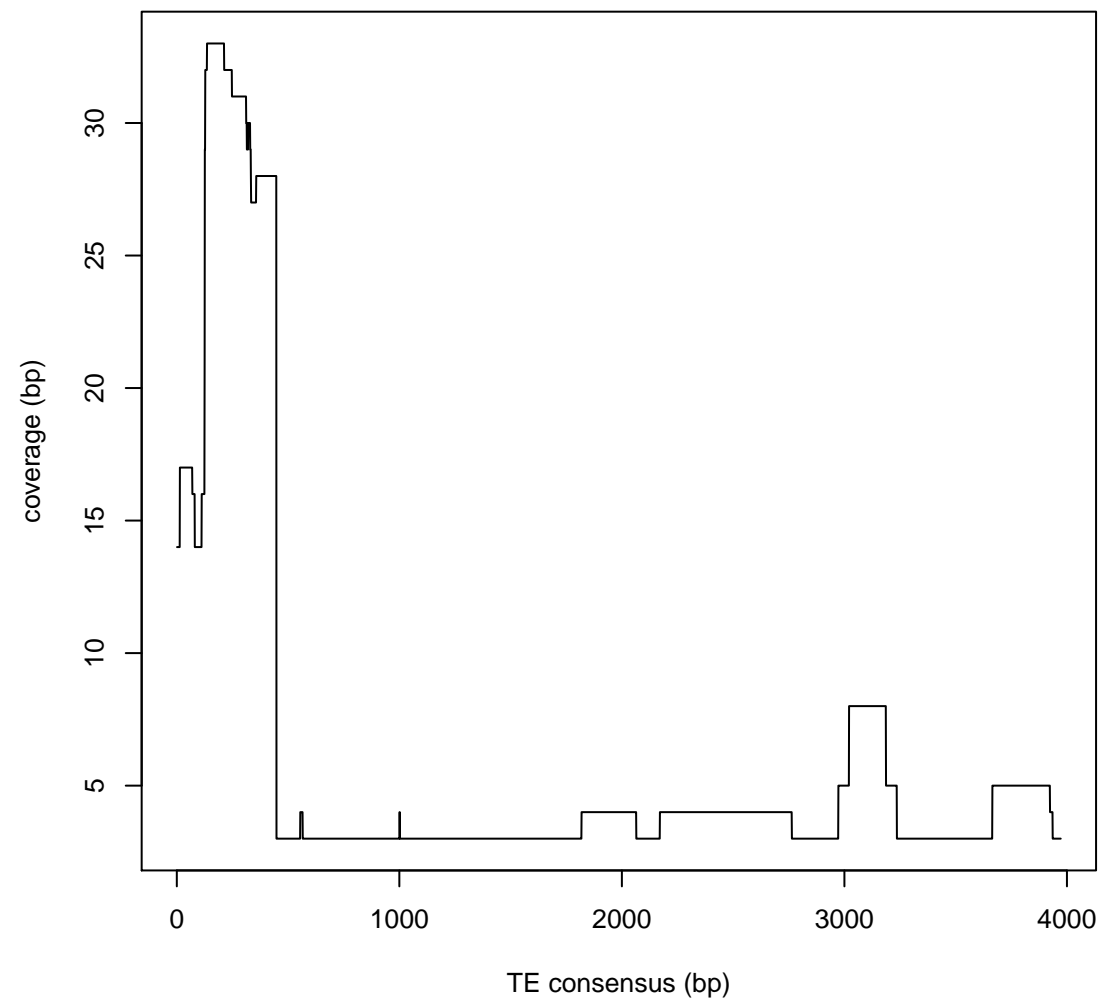


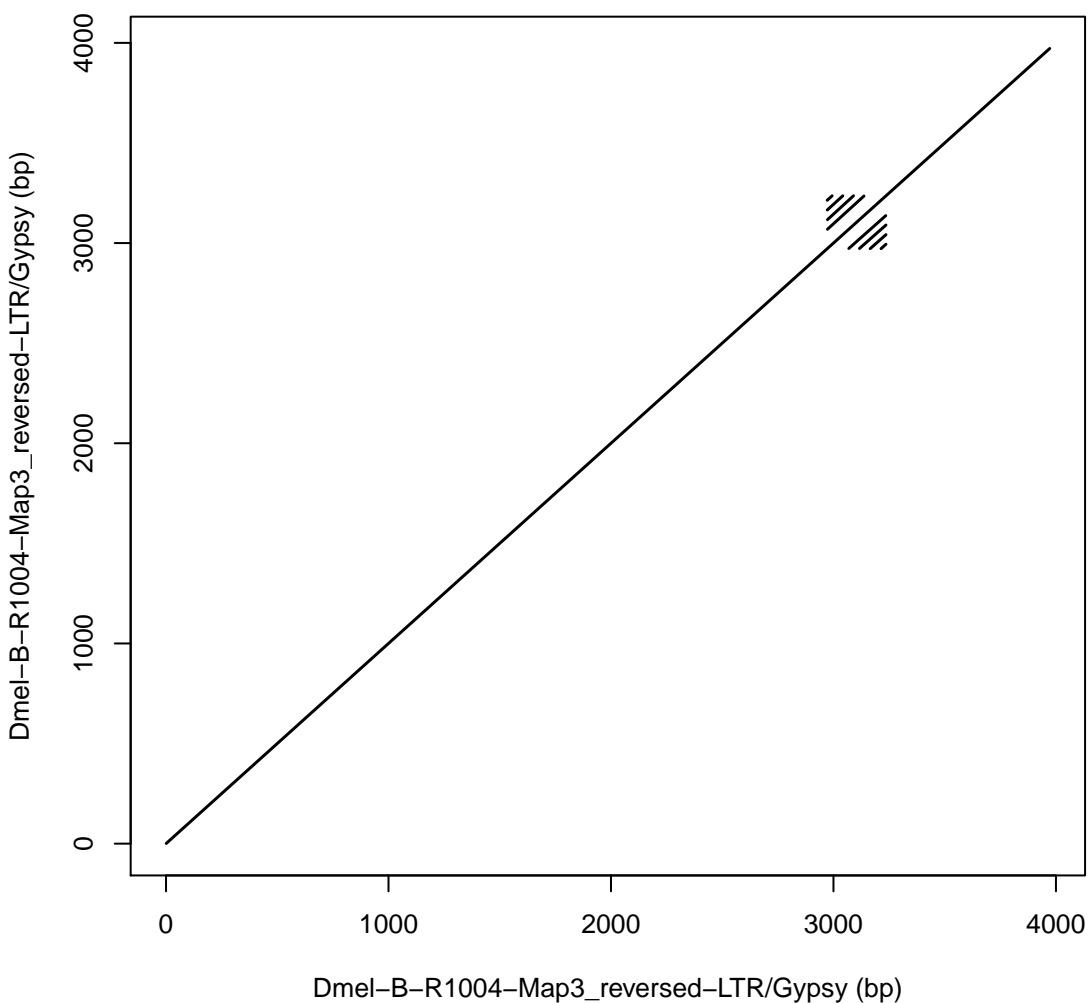
TE: Dmel-B-R1004-Map3_reversed-LTR/Gypsy
consensus size: 3972bp; fragments: 49; full length: 2 (>=3574.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

