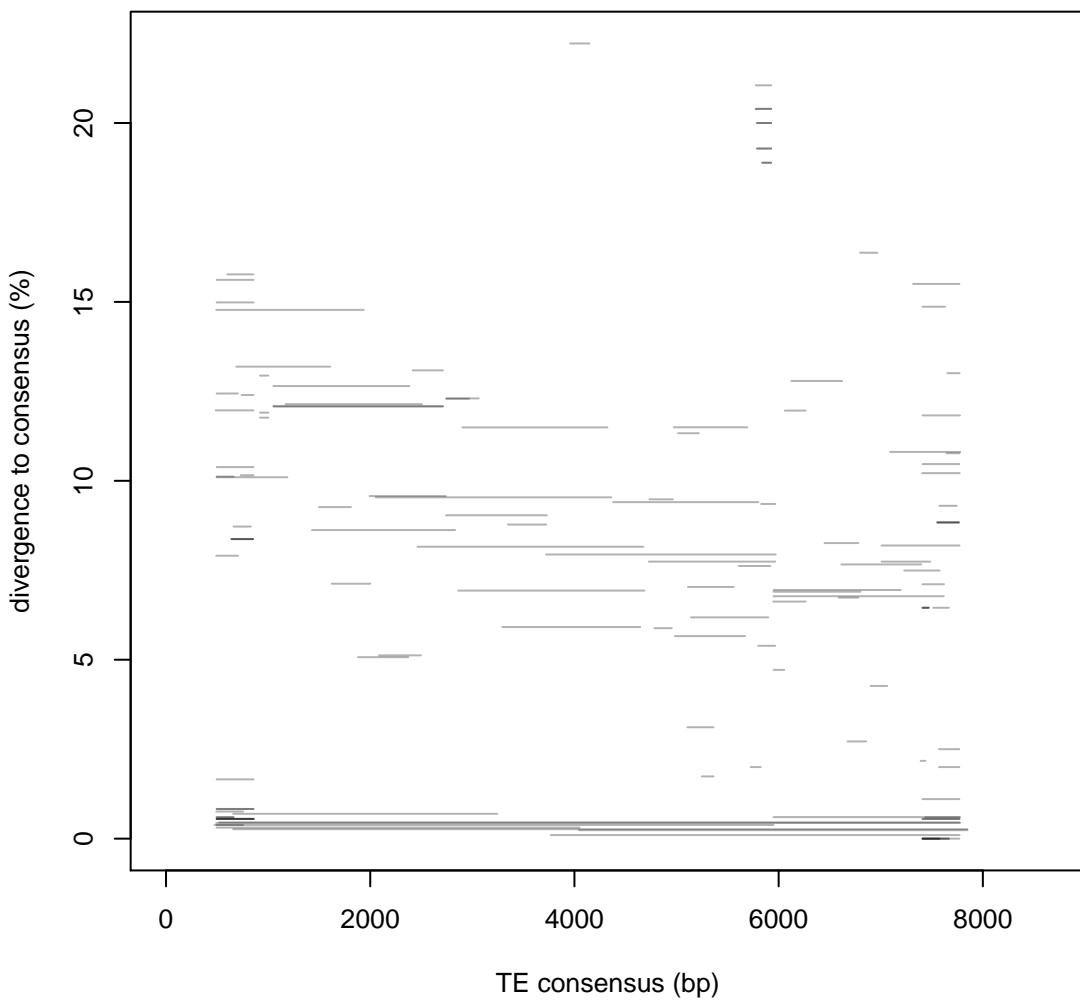
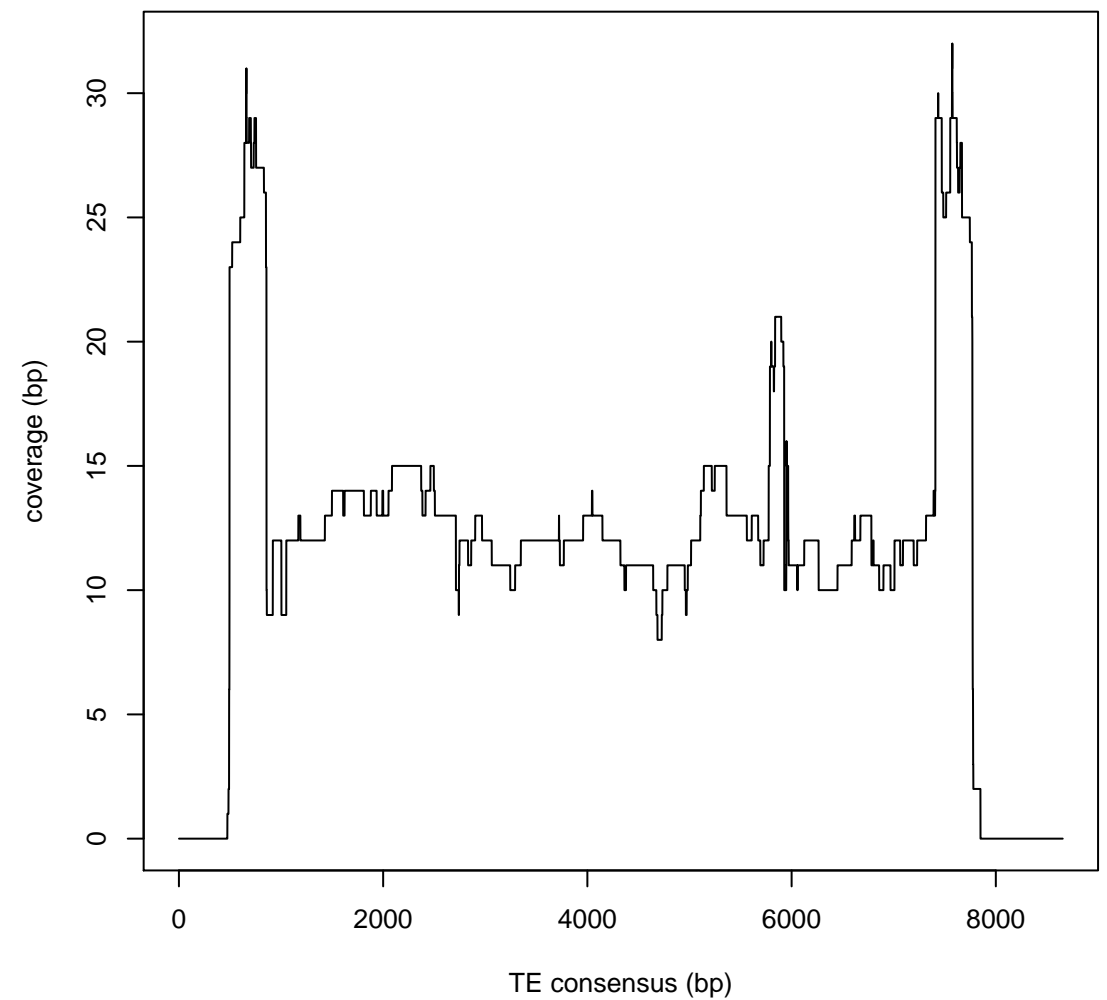


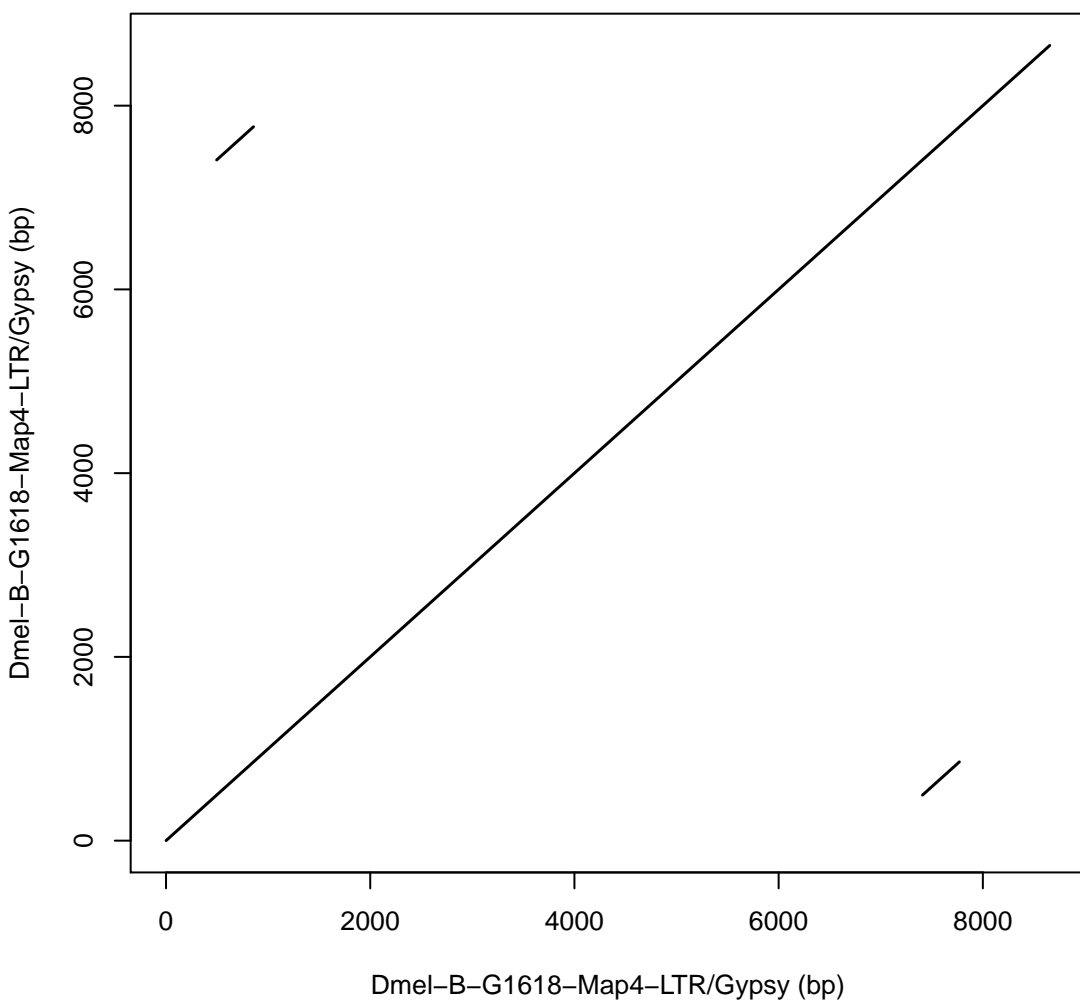
TE: Dmel-B-G1618-Map4-LTR/Gypsy
consensus size: 8655bp; fragments: 131; full length: 0 (>=7789.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

