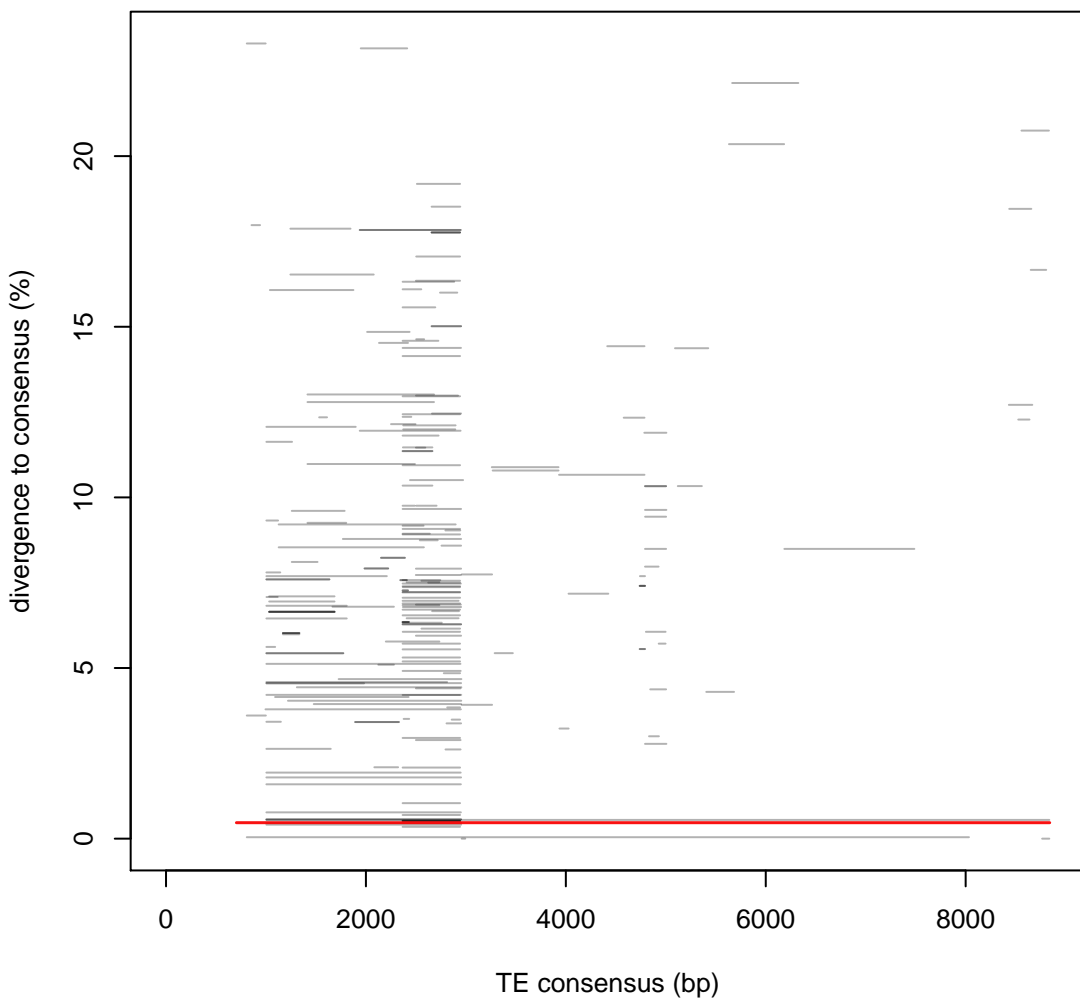
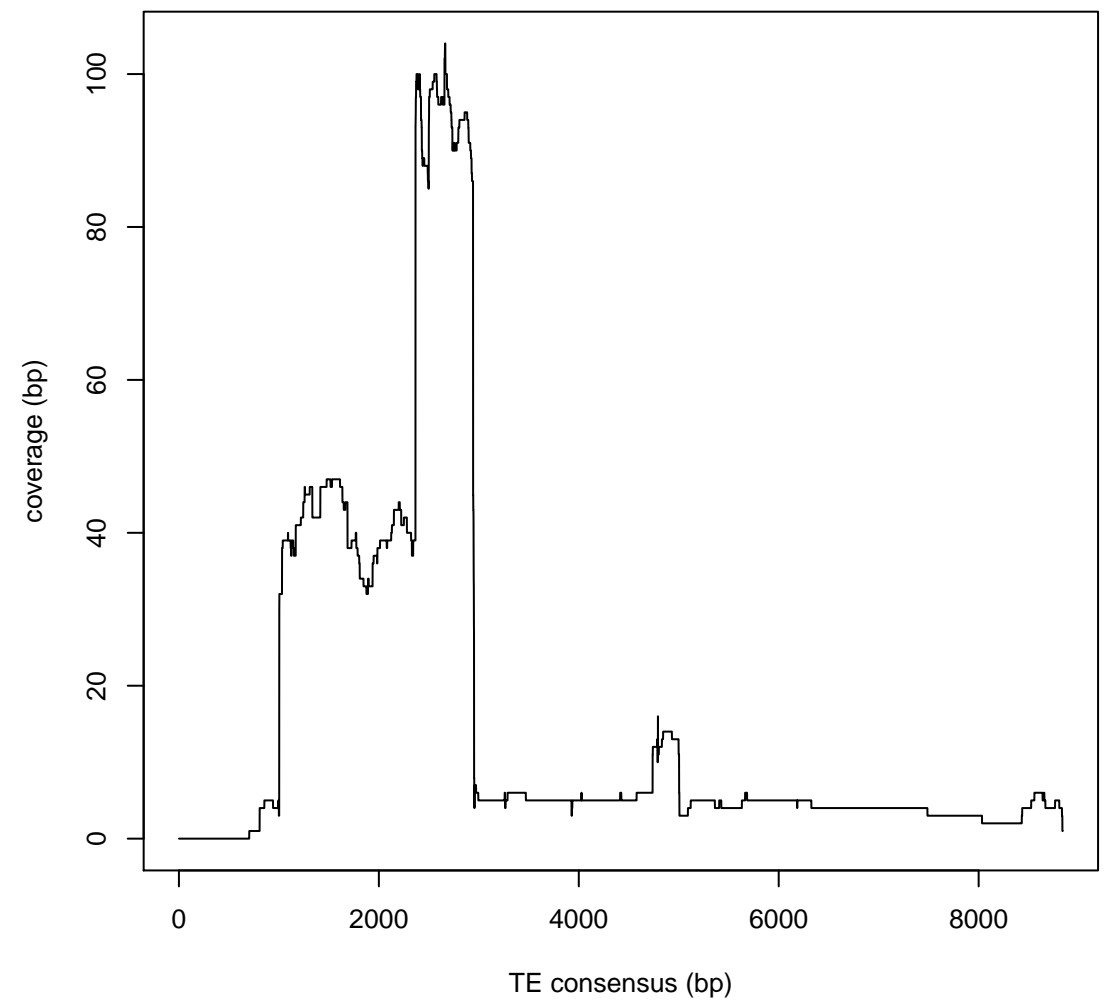


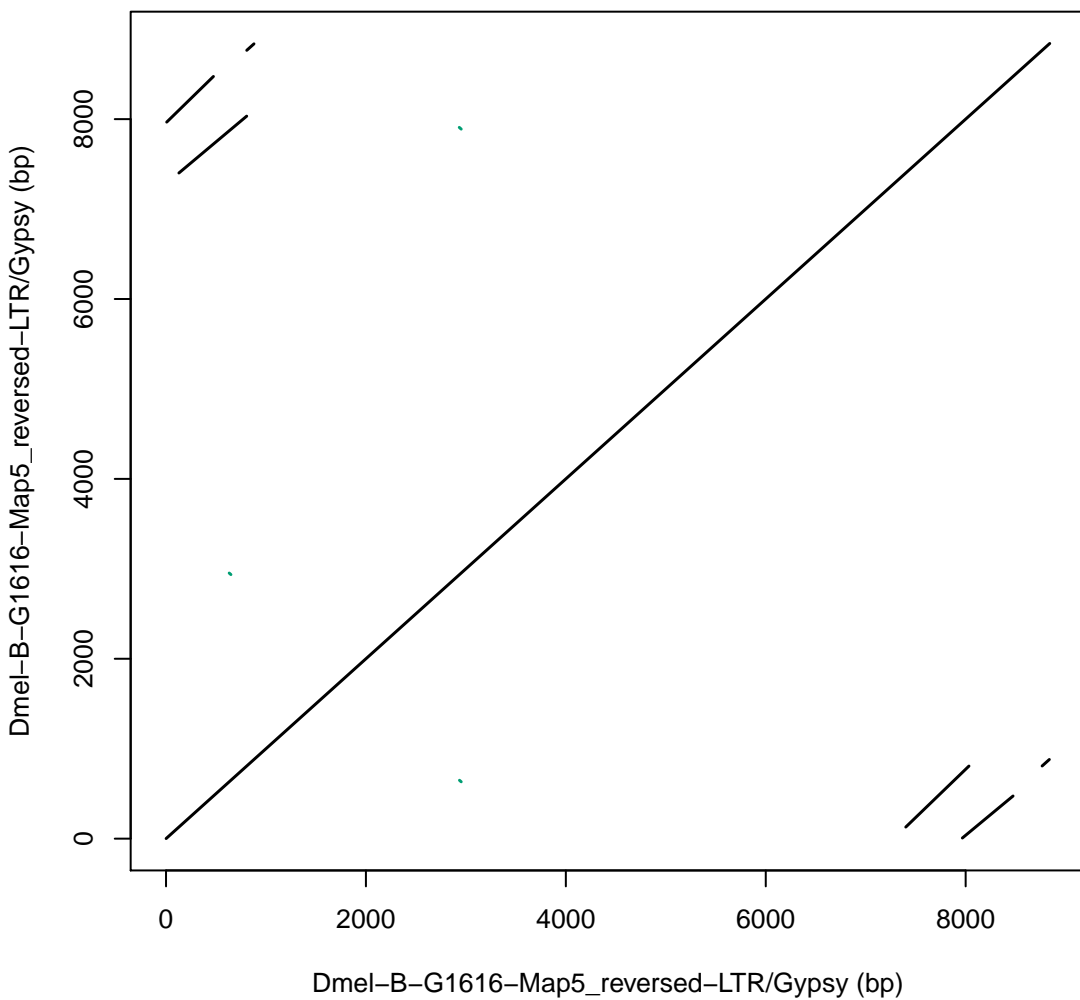
TE: Dmel-B-G1616-Map5_reversed-LTR/Gypsy
consensus size: 8841bp; fragments: 220; full length: 1 (>=7956.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

