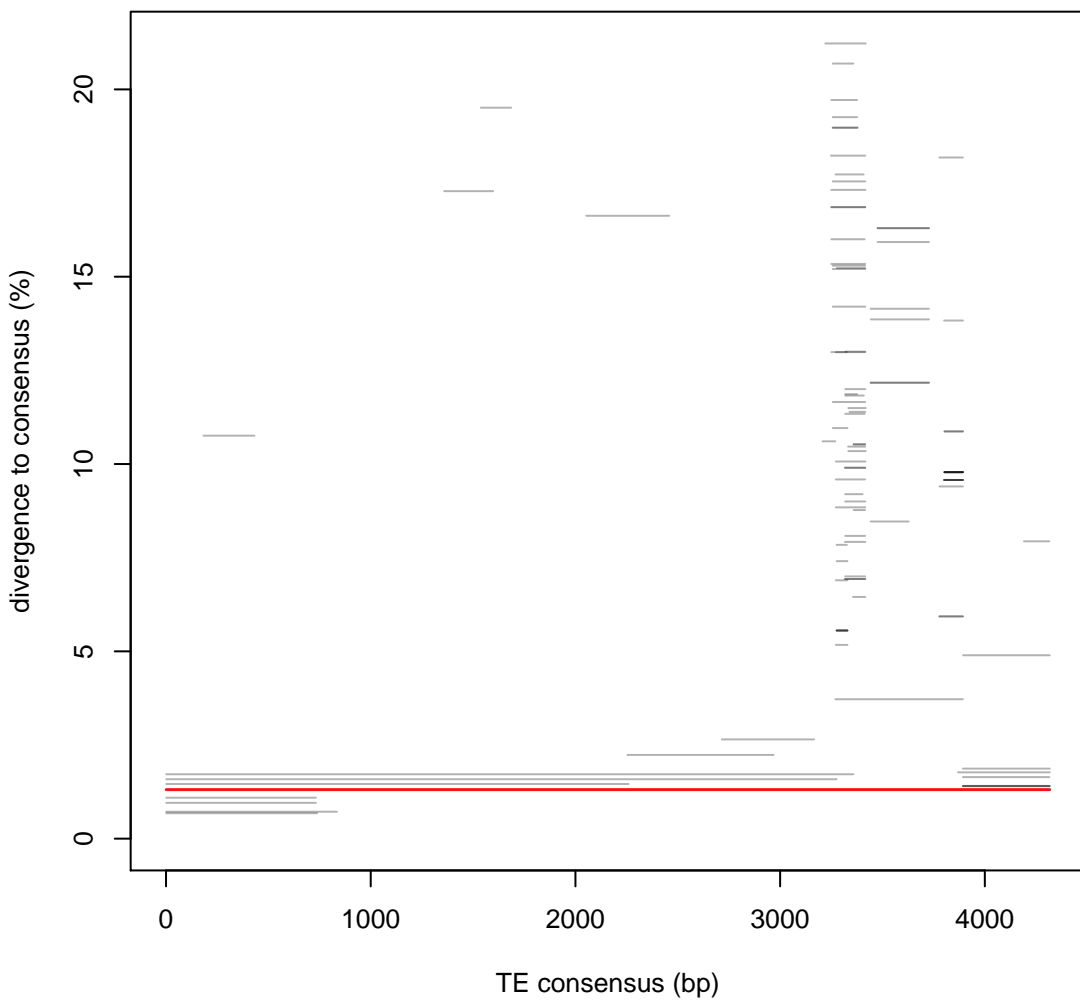
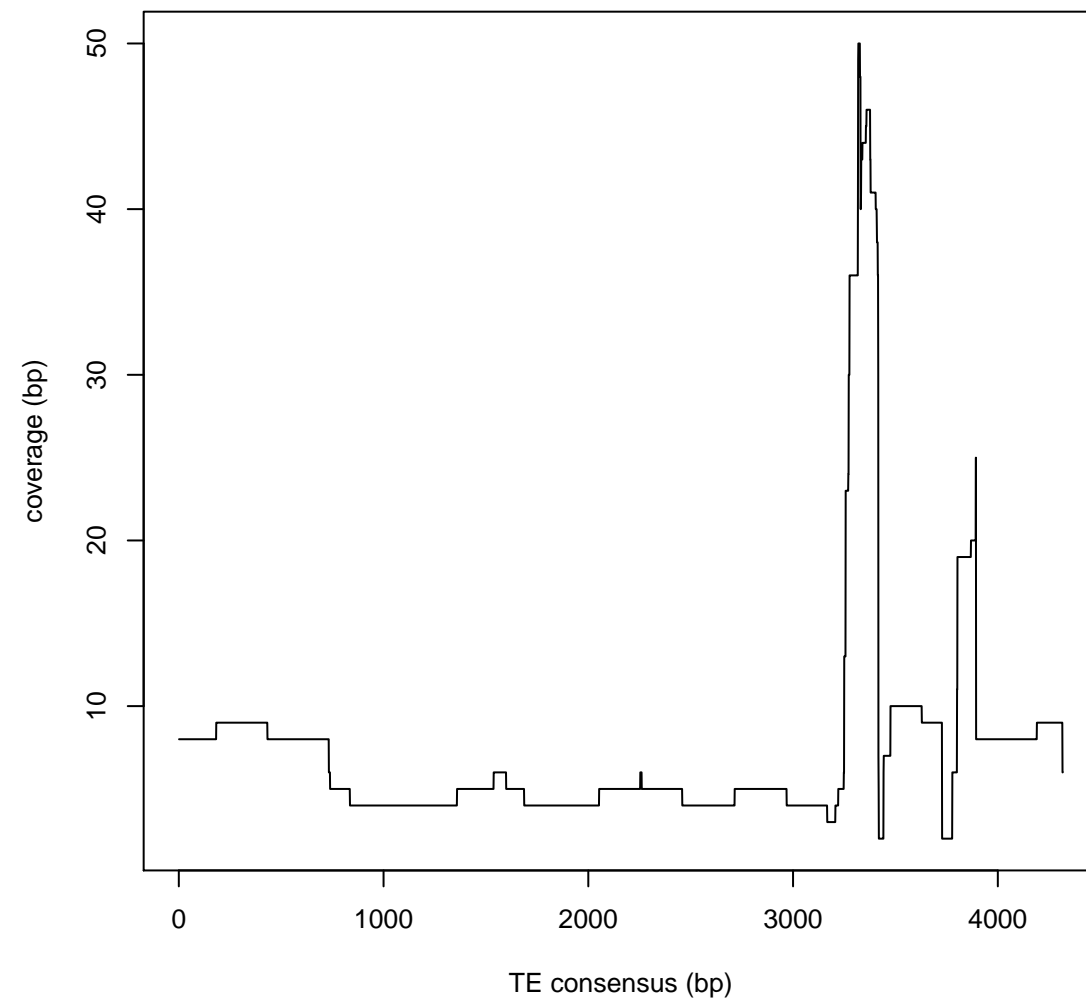


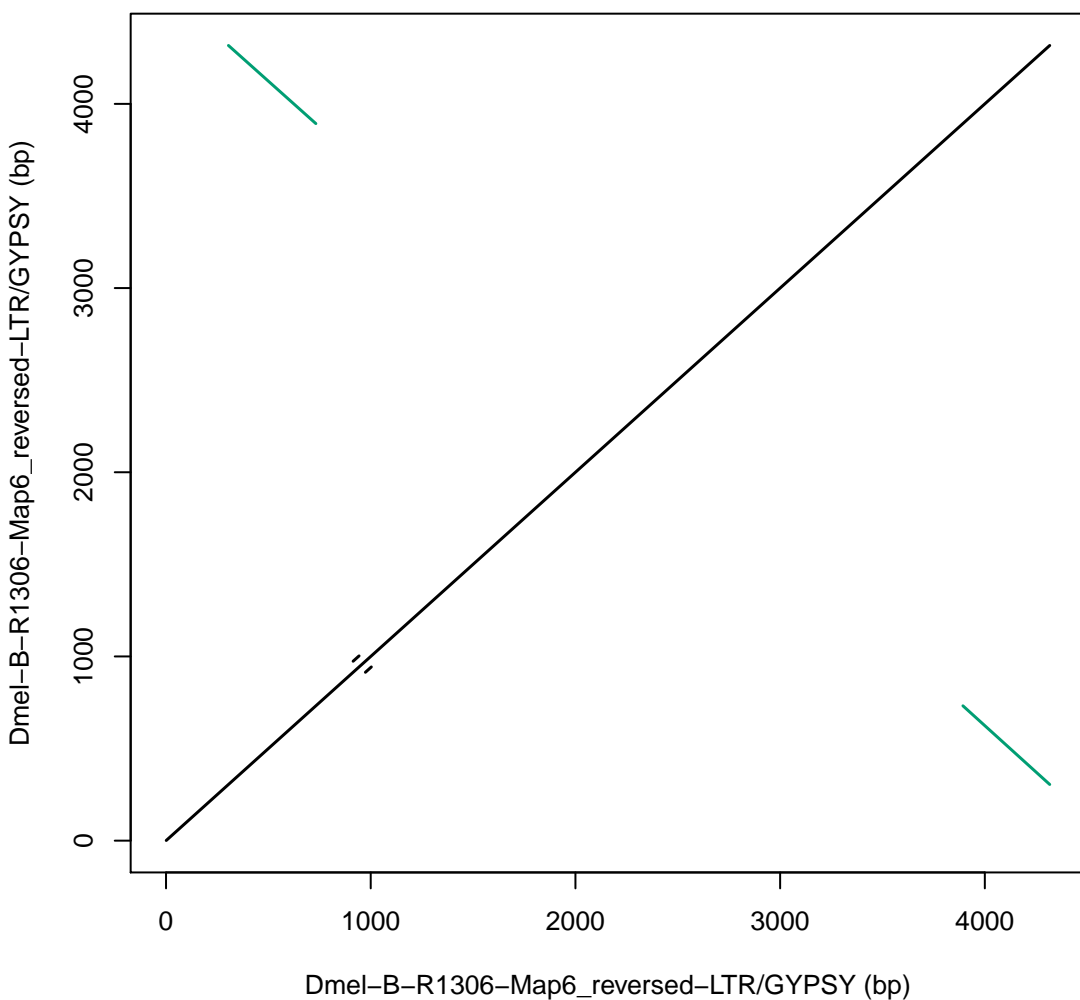
TE: Dmel-B-R1306-Map6\_reversed-LTR/GYPY  
consensus size: 4317bp; fragments: 104; full length: 1 (>=3885.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

