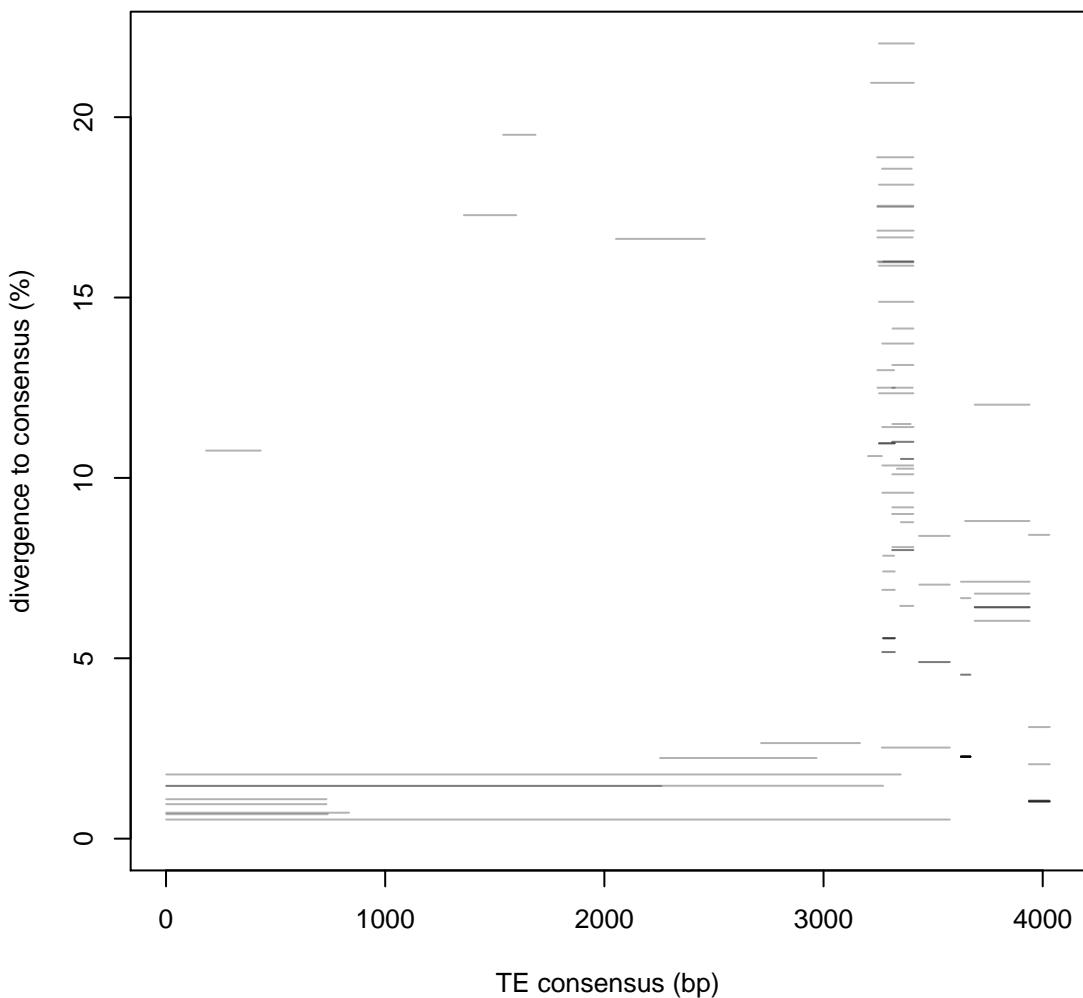
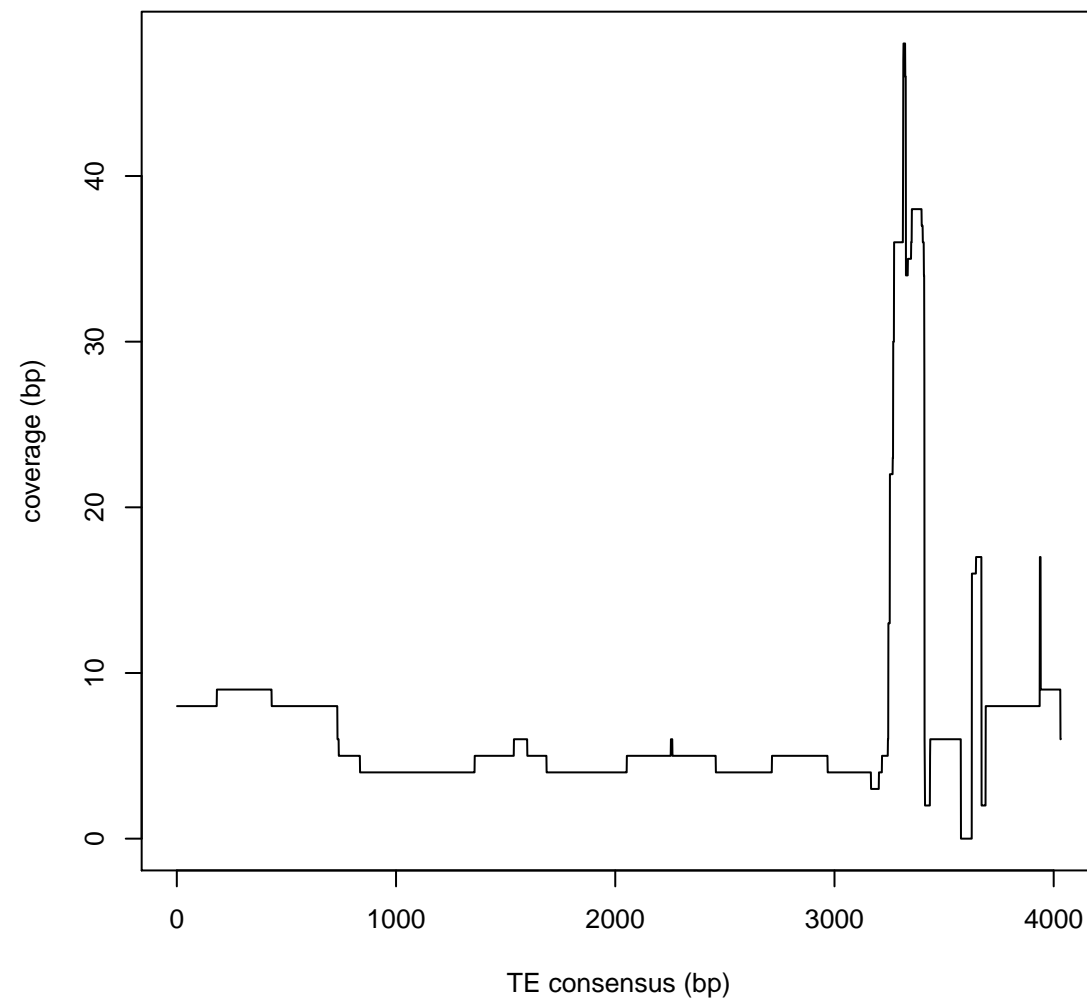


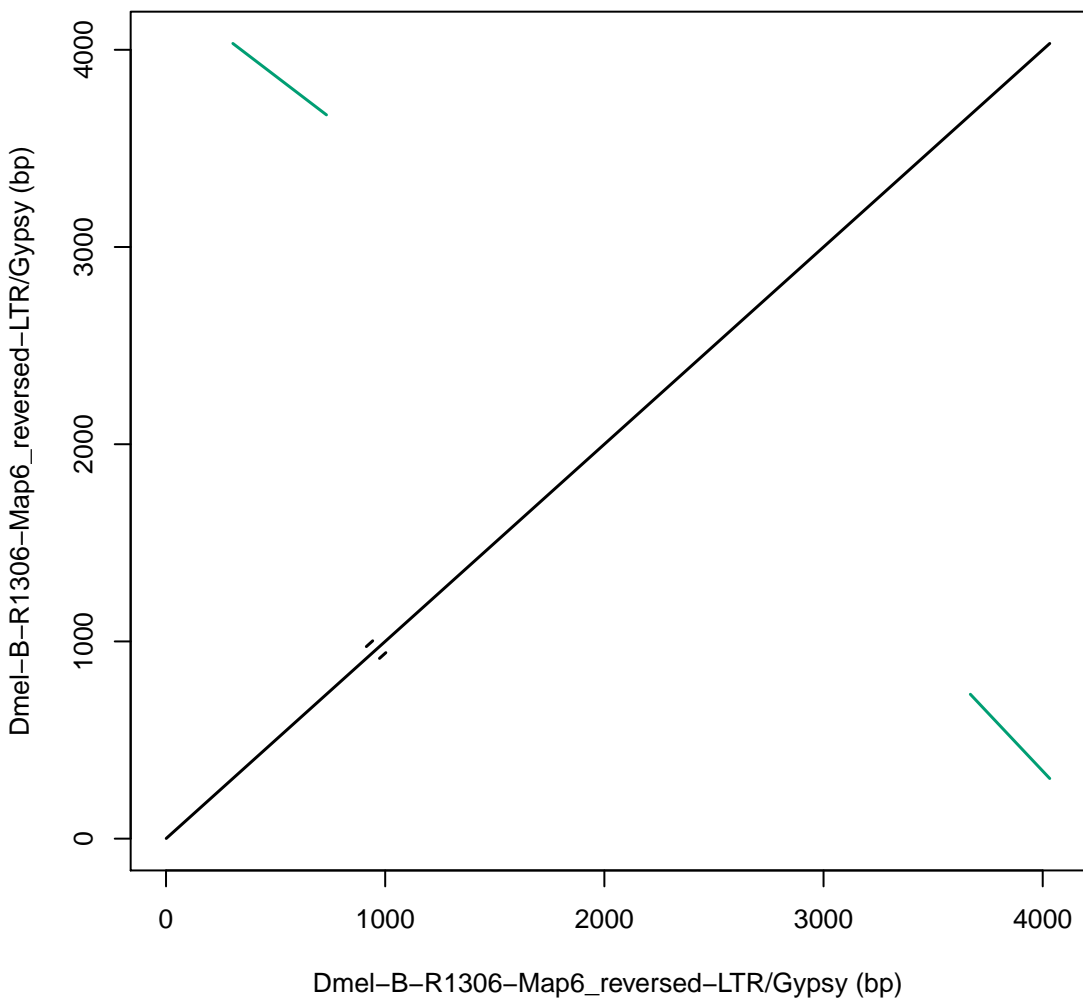
TE: Dmel-B-R1306-Map6_reversed-LTR/Gypsy
consensus size: 4032bp; fragments: 102; full length: 0 (>=3628.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

