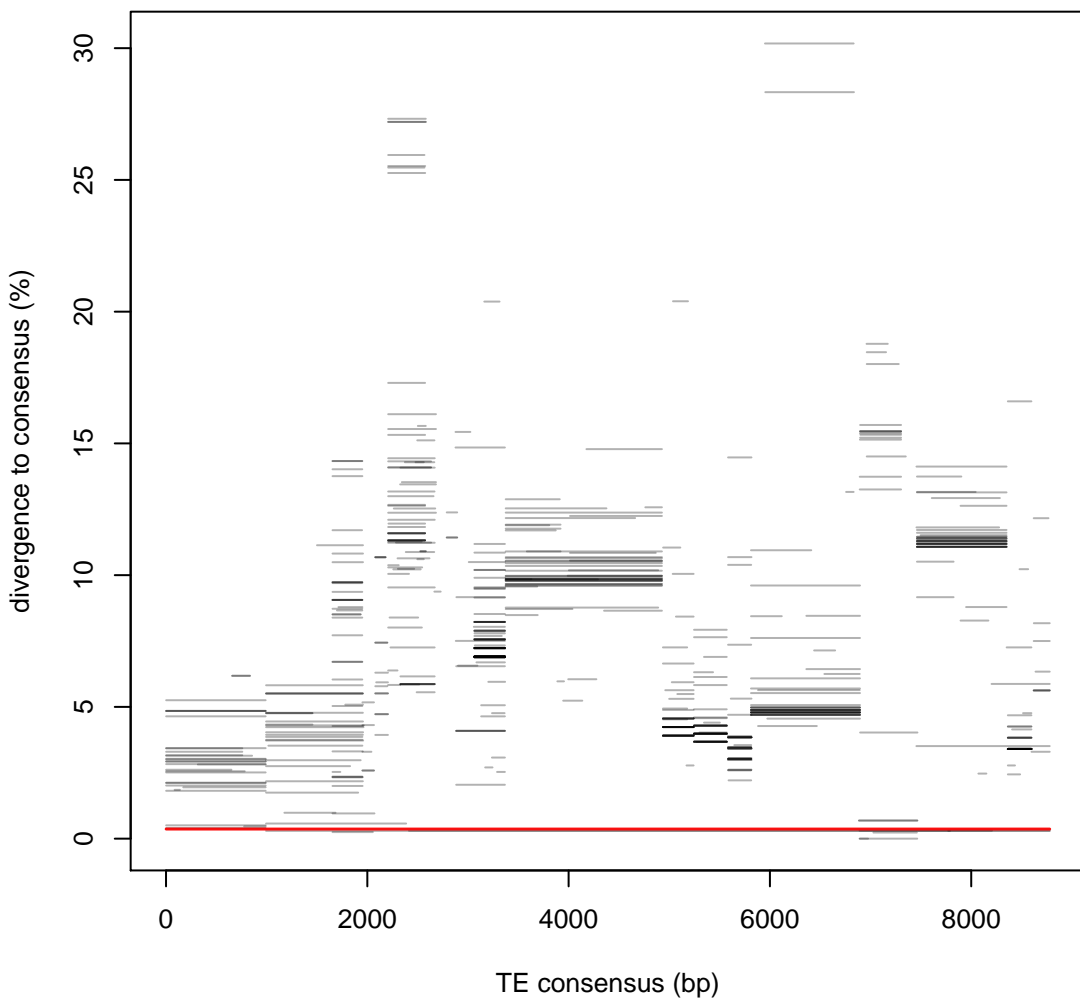
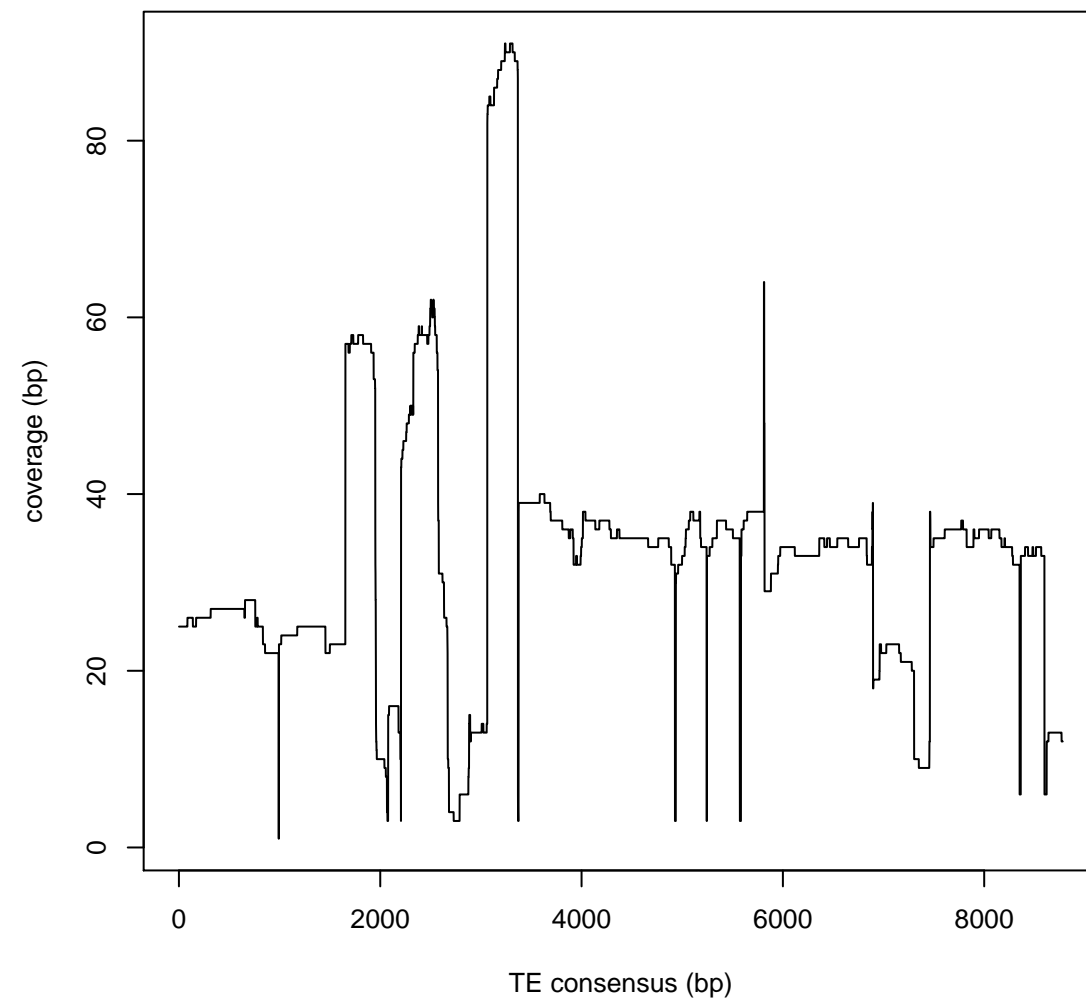


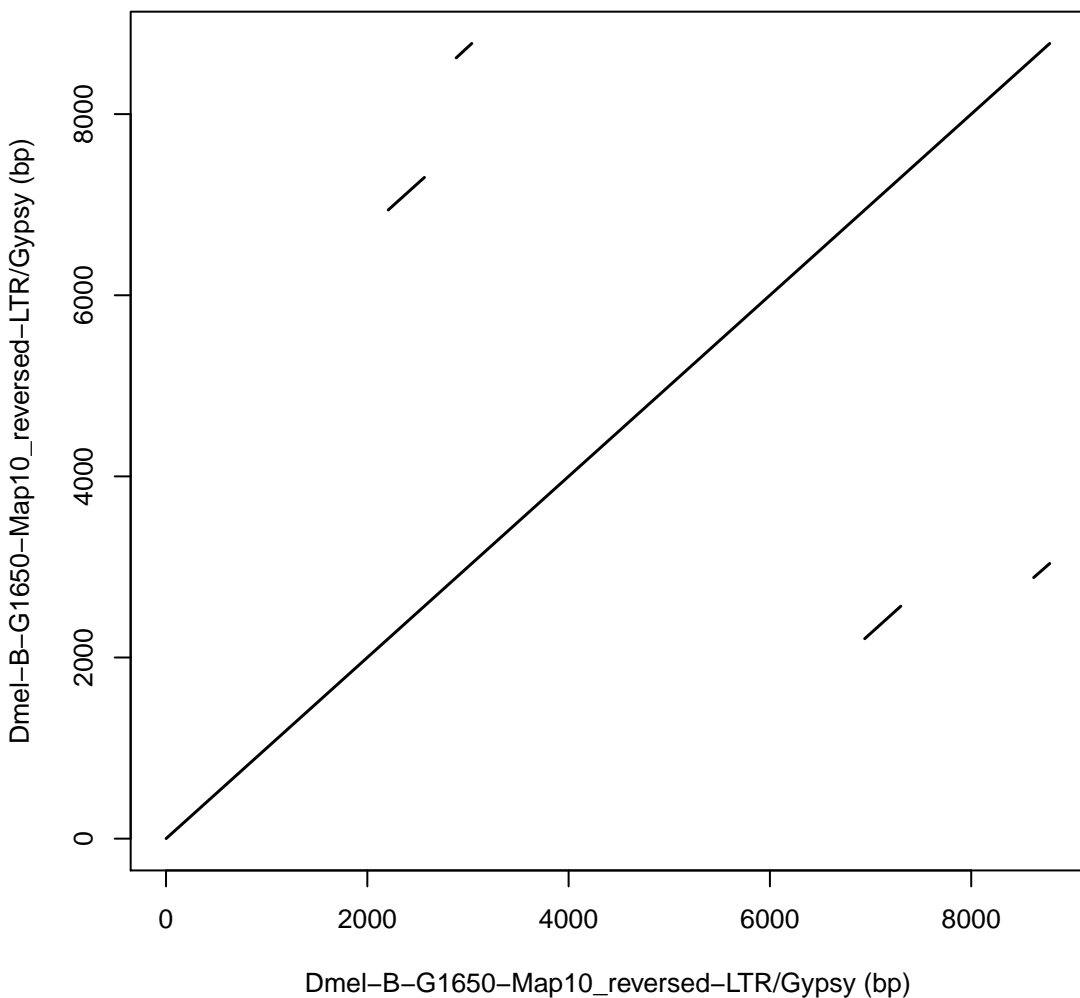
TE: Dmel-B-G1650-Map10\_reversed-LTR/Gypsy  
consensus size: 8780bp; fragments: 560; full length: 1 (>=7902bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

