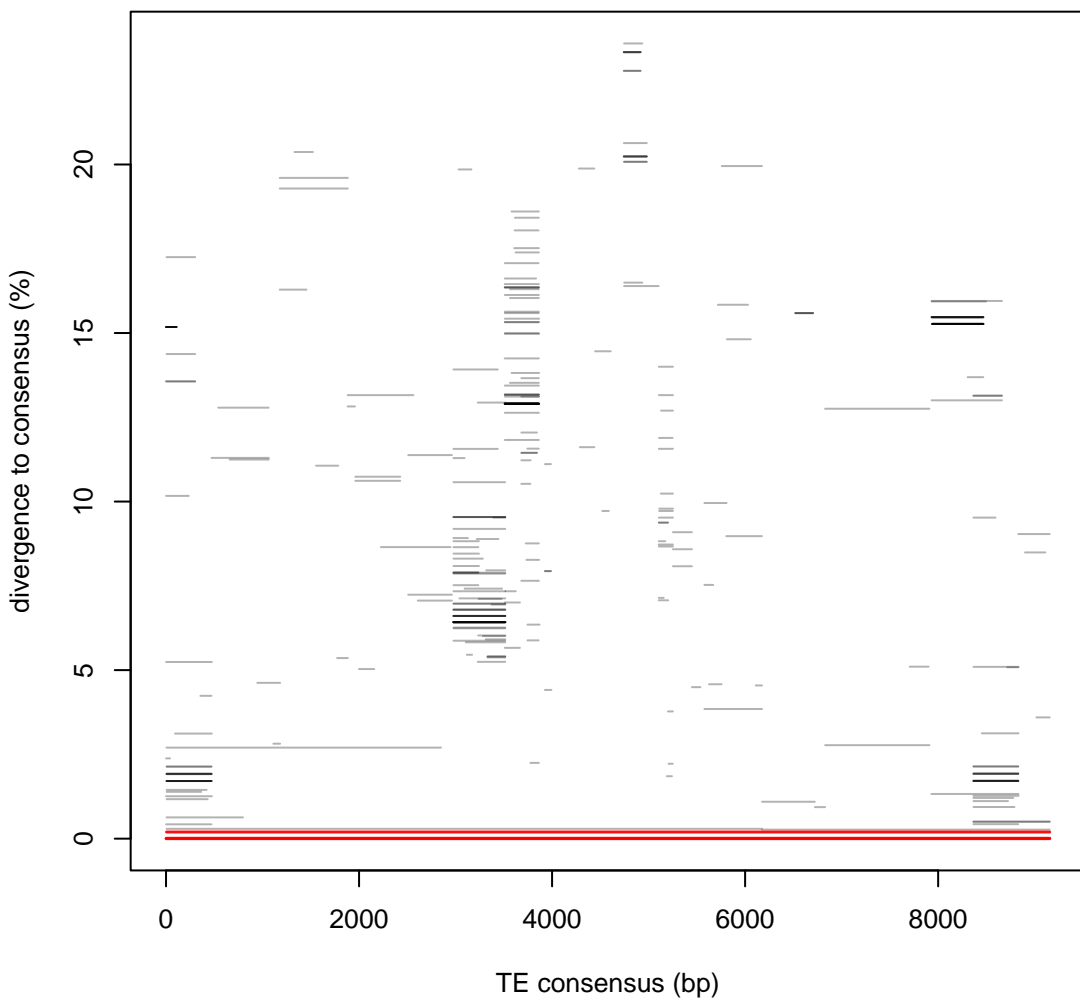
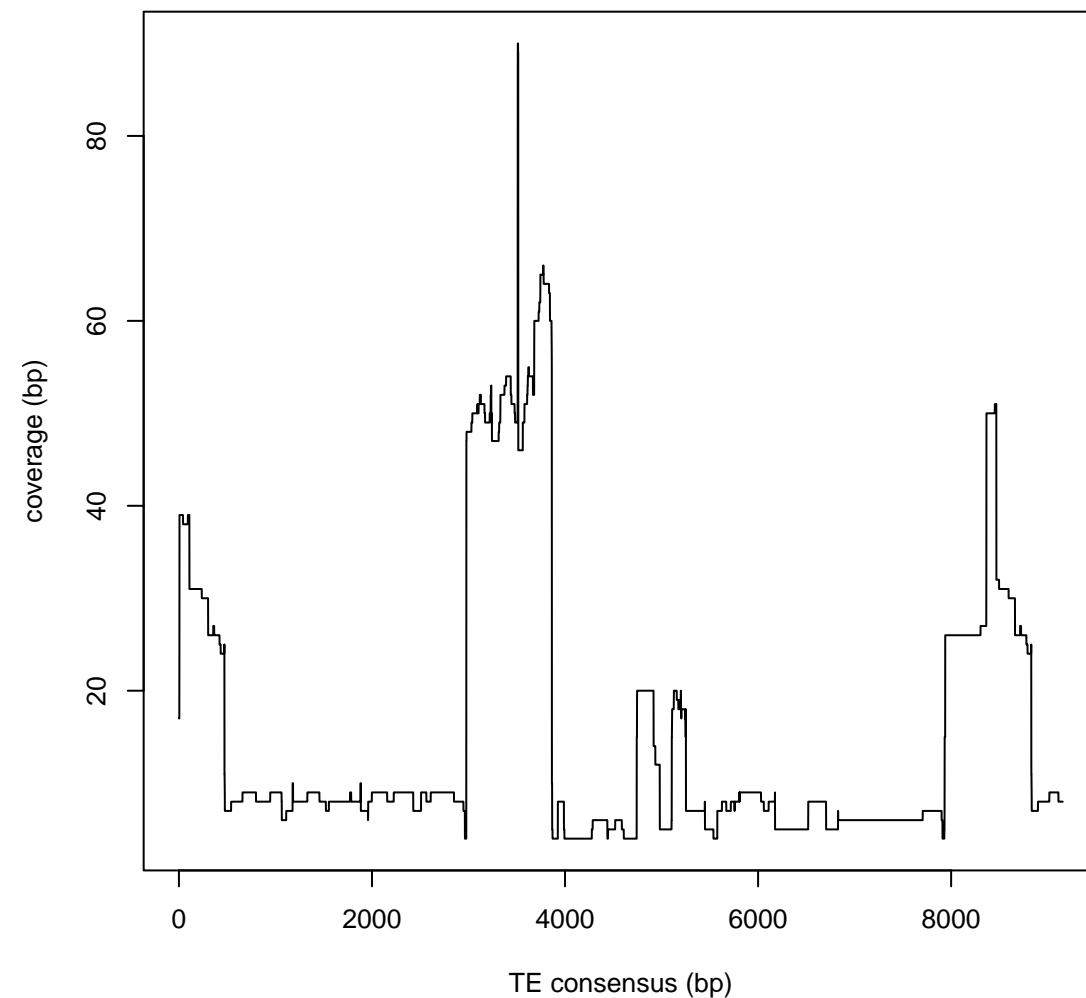


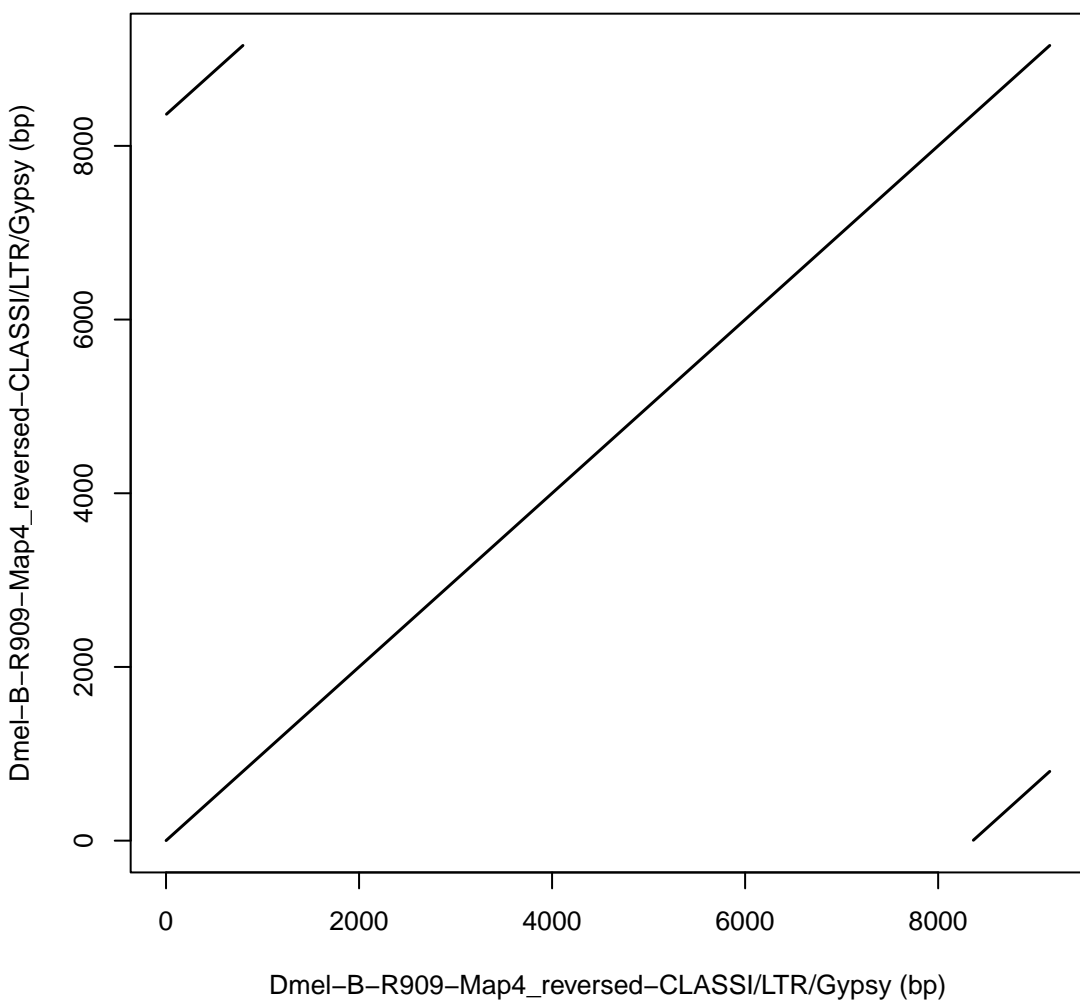
TE: Dmel-B-R909-Map4\_reversed-CLASSI/LTR/Gypsy  
consensus size: 9156bp; fragments: 304; full length: 3 (>=8240.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

