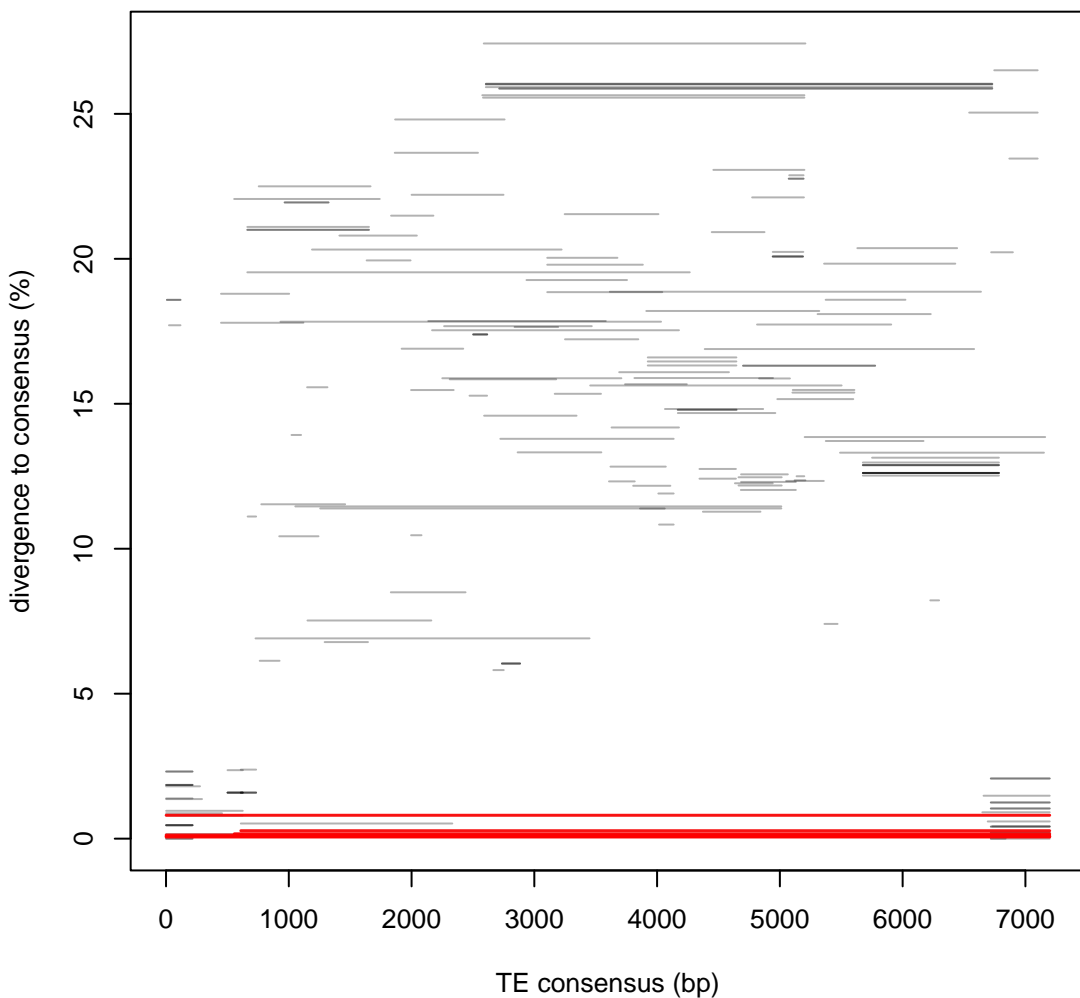
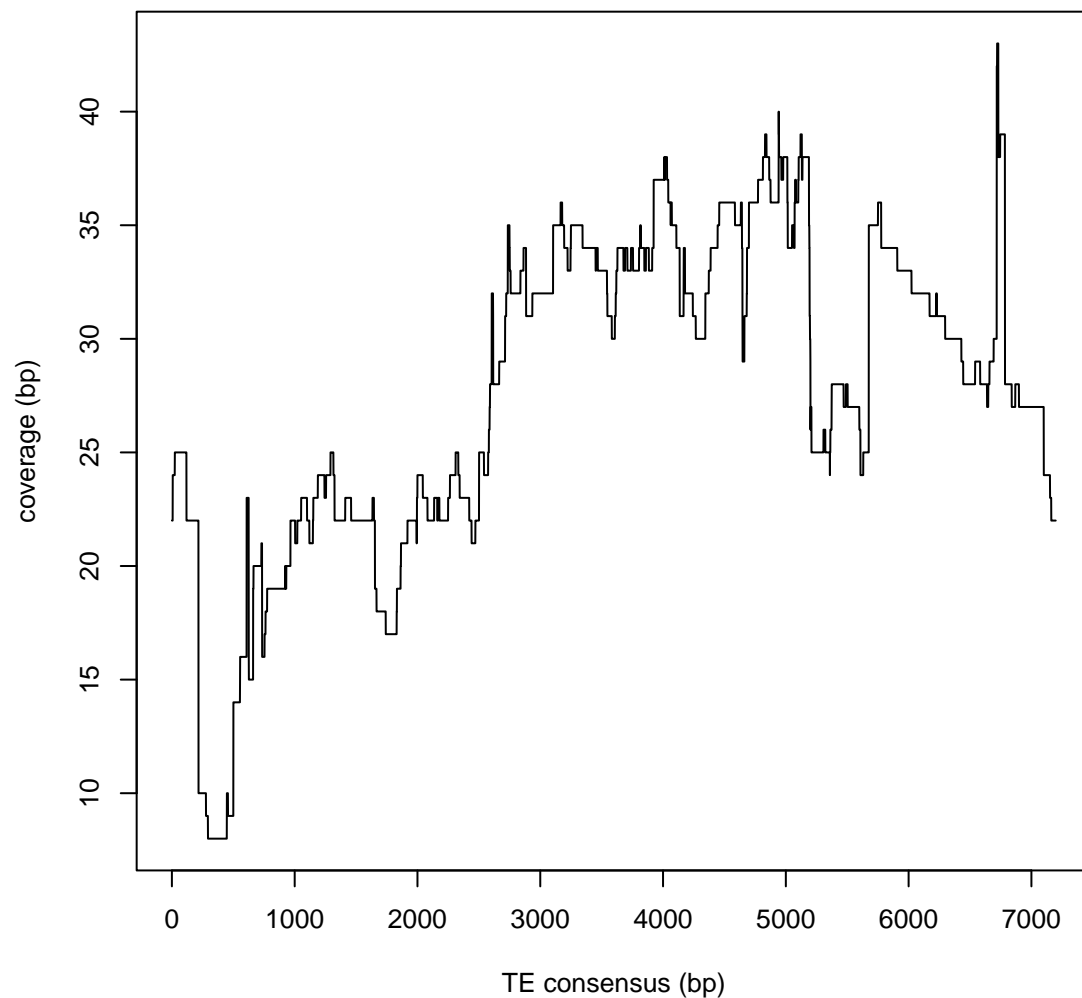


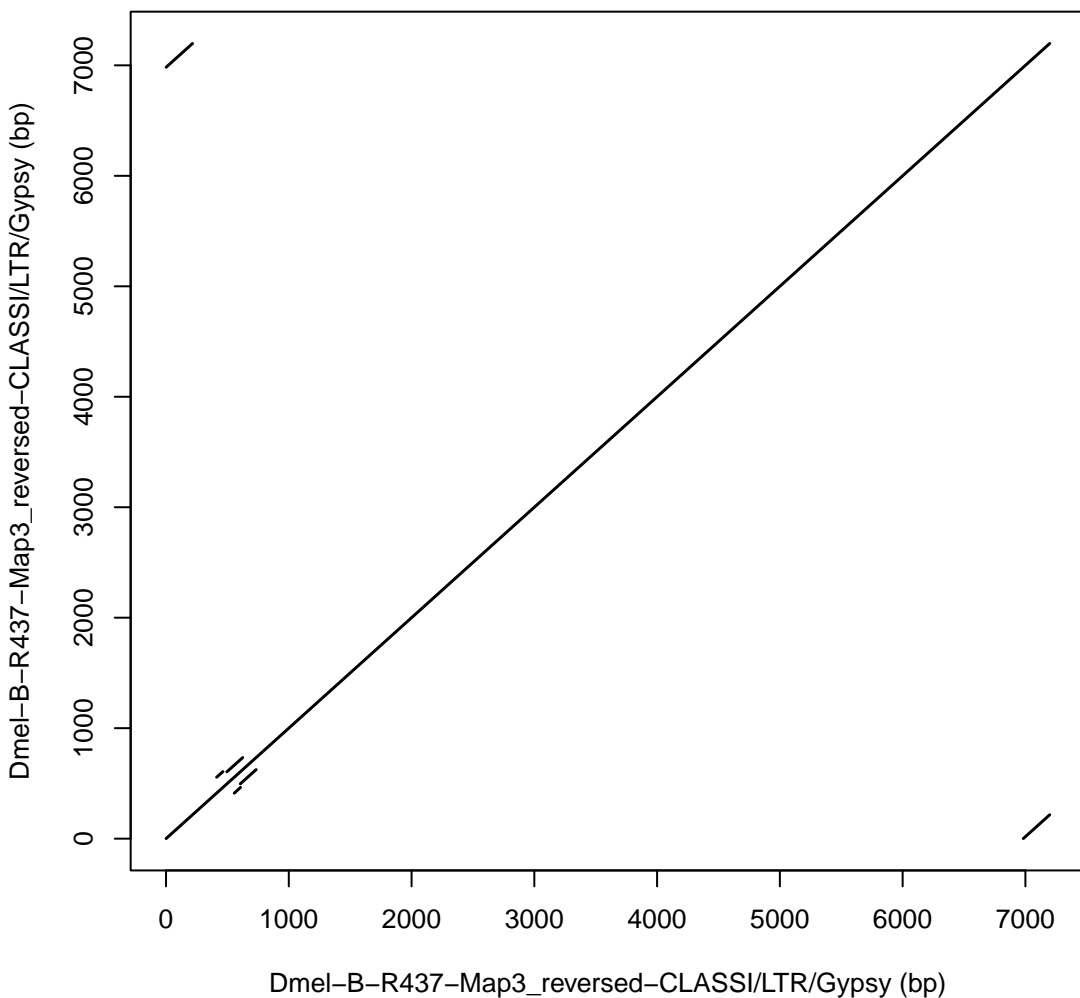
TE: Dmel-B-R437-Map3\_reversed-CLASSI/LTR/Gypsy  
consensus size: 7198bp; fragments: 194; full length: 7 (>=6478.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

