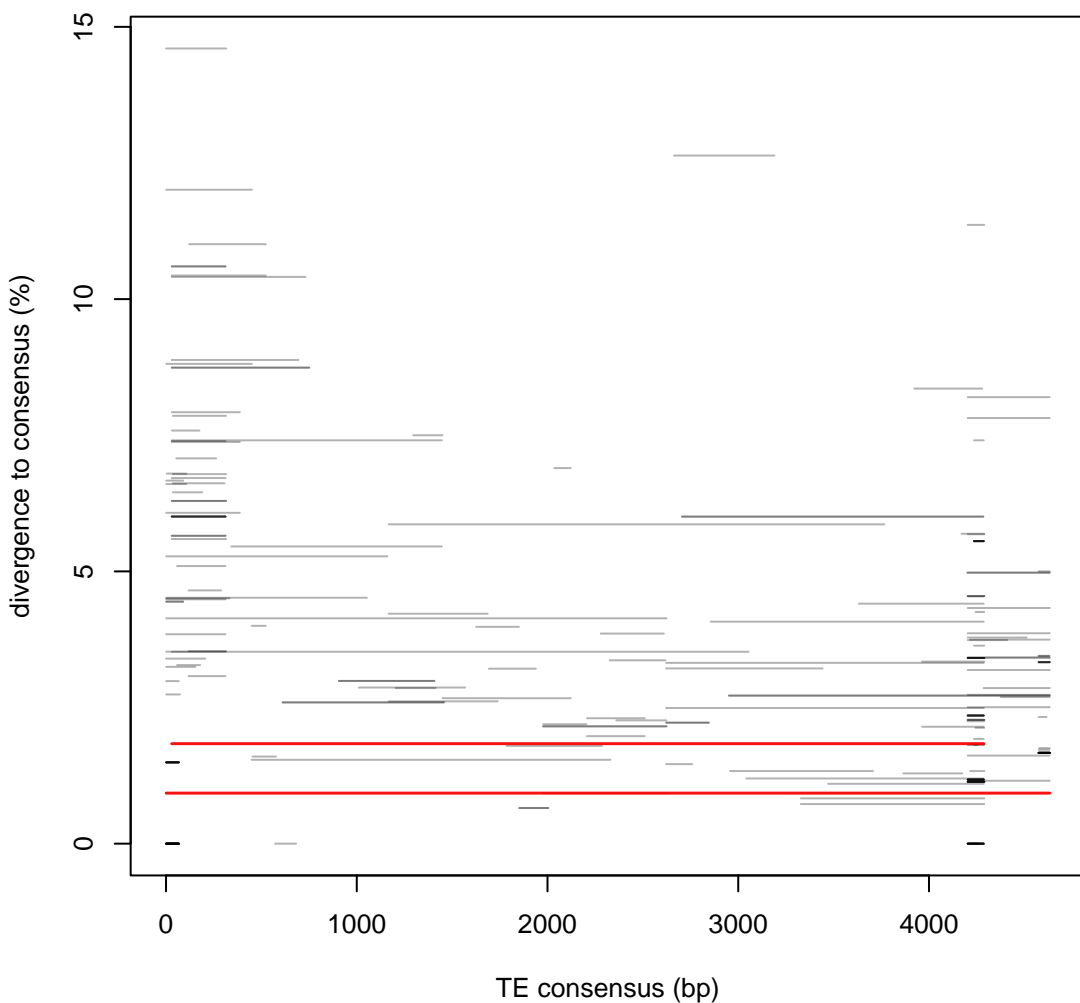
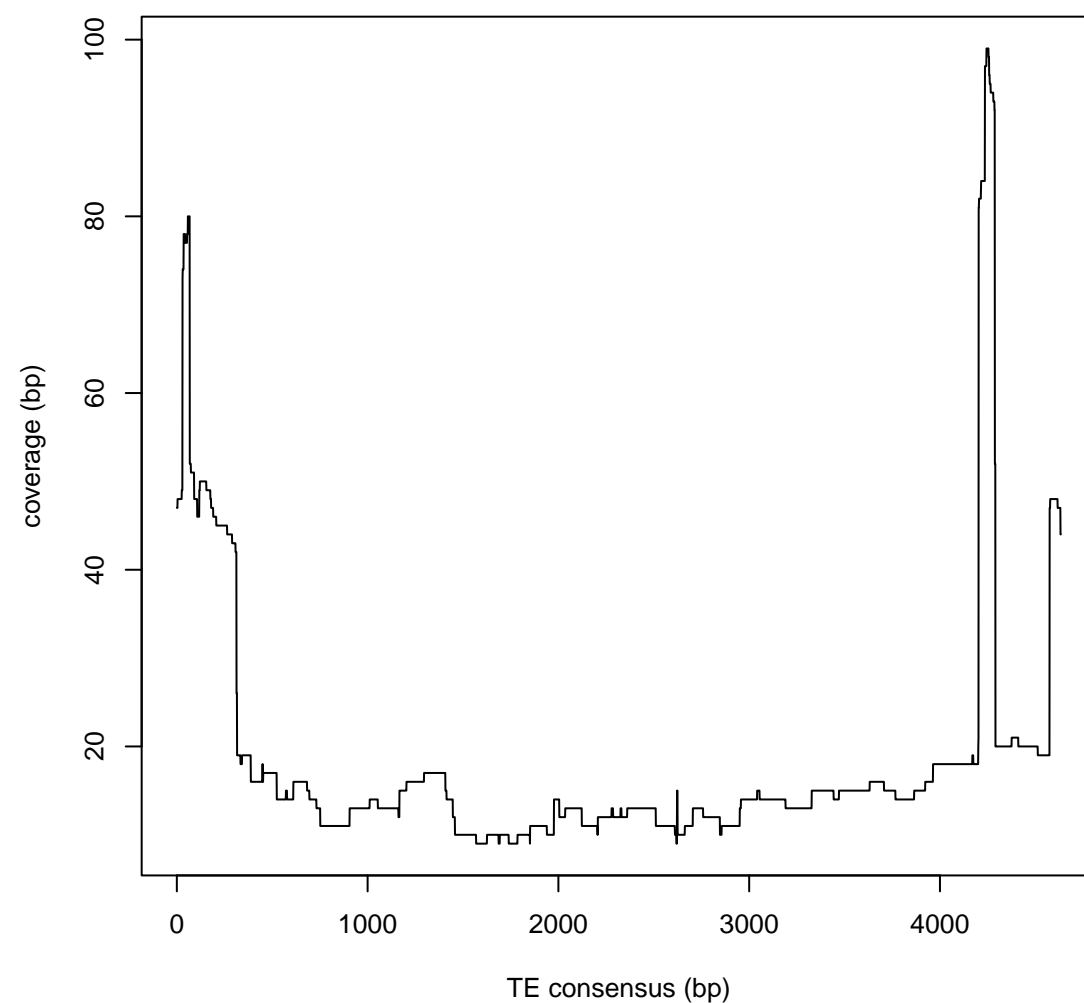


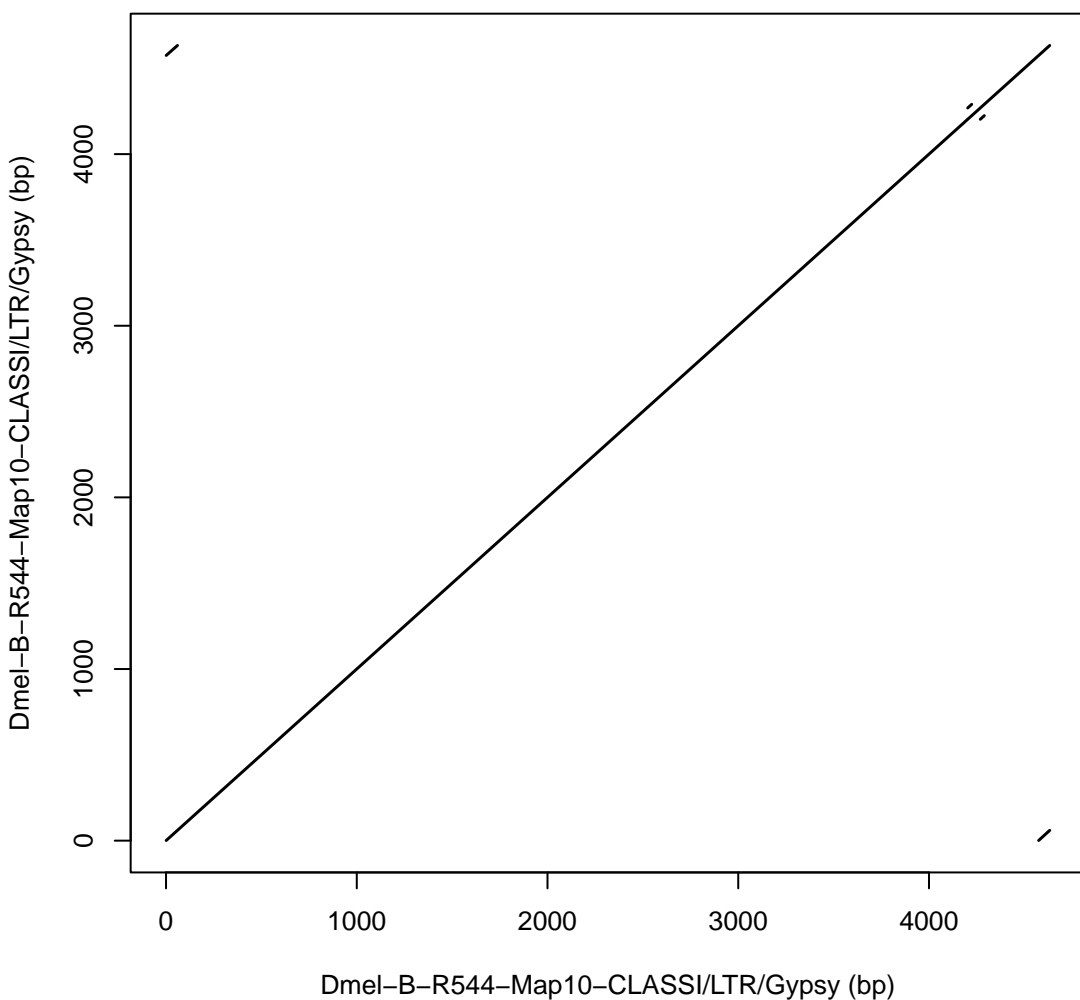
TE: Dmel-B-R544-Map10-CLASSI/LTR/Gypsy  
consensus size: 4633bp; fragments: 251; full length: 2 (>=4169.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

