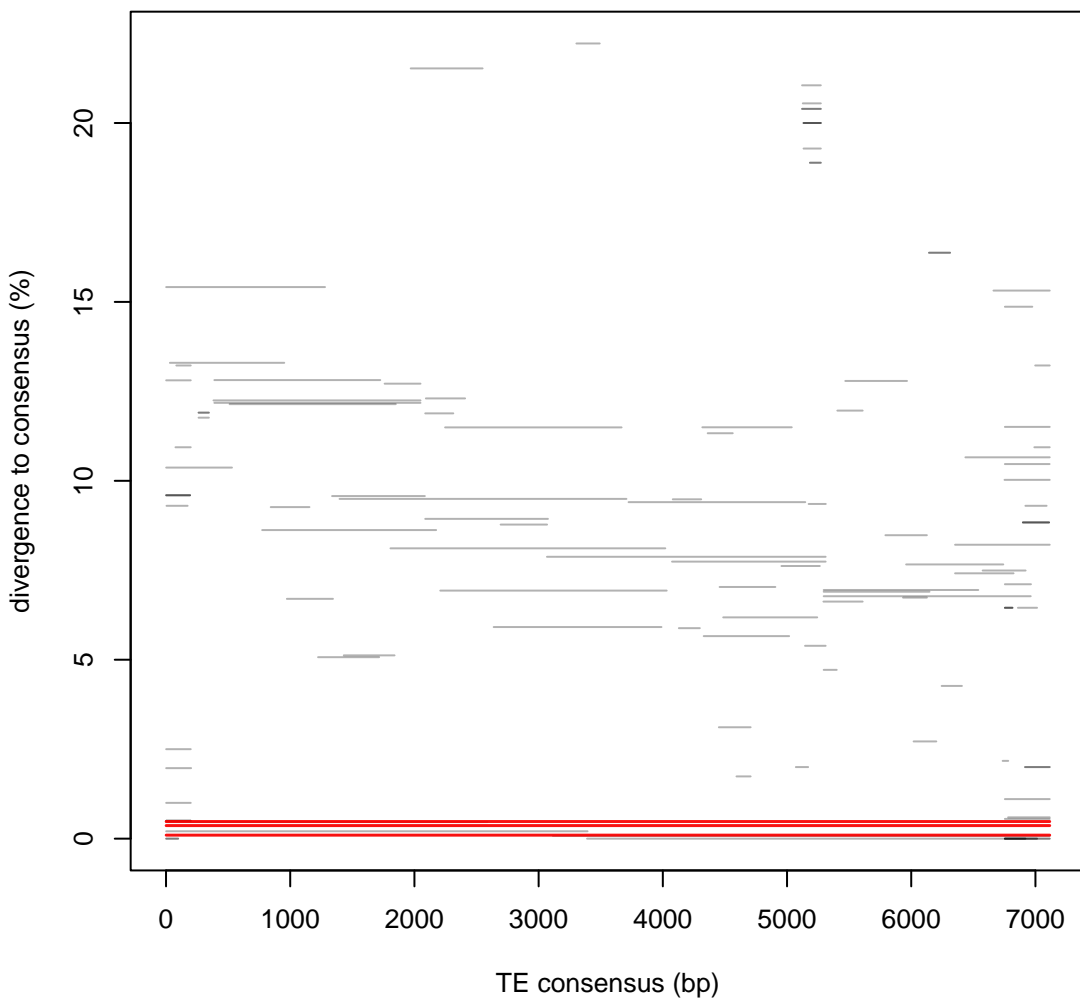
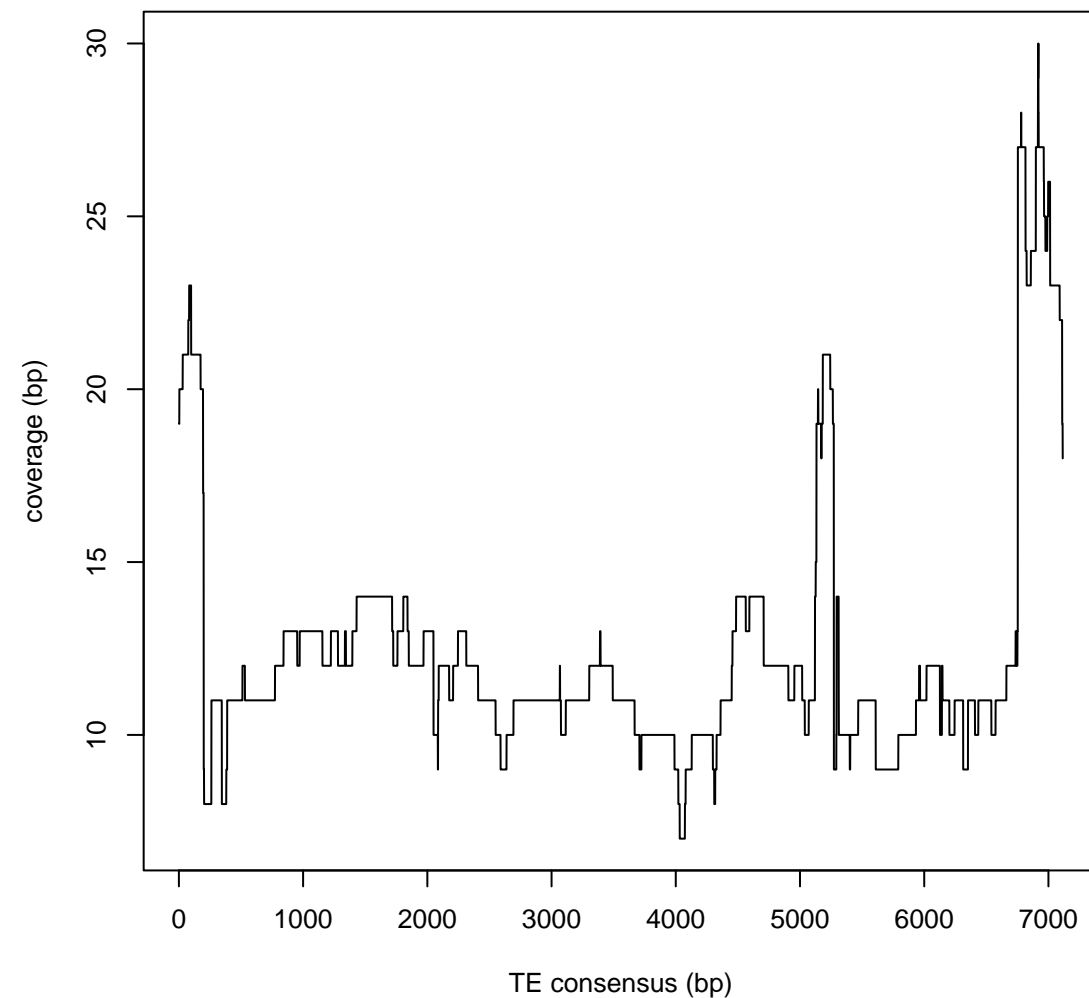


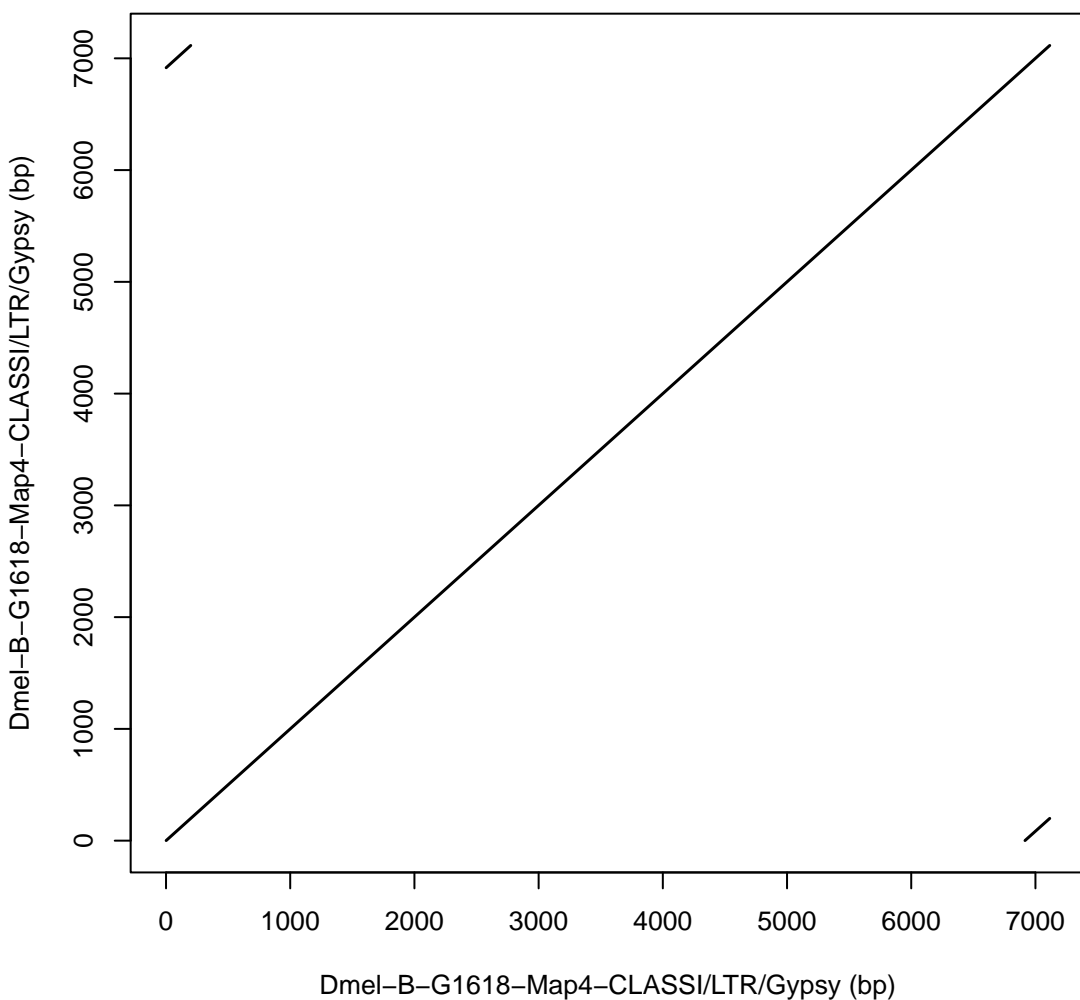
TE: Dmel-B-G1618-Map4-CLASSI/LTR/Gypsy  
consensus size: 7115bp; fragments: 121; full length: 3 (>=6403.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

