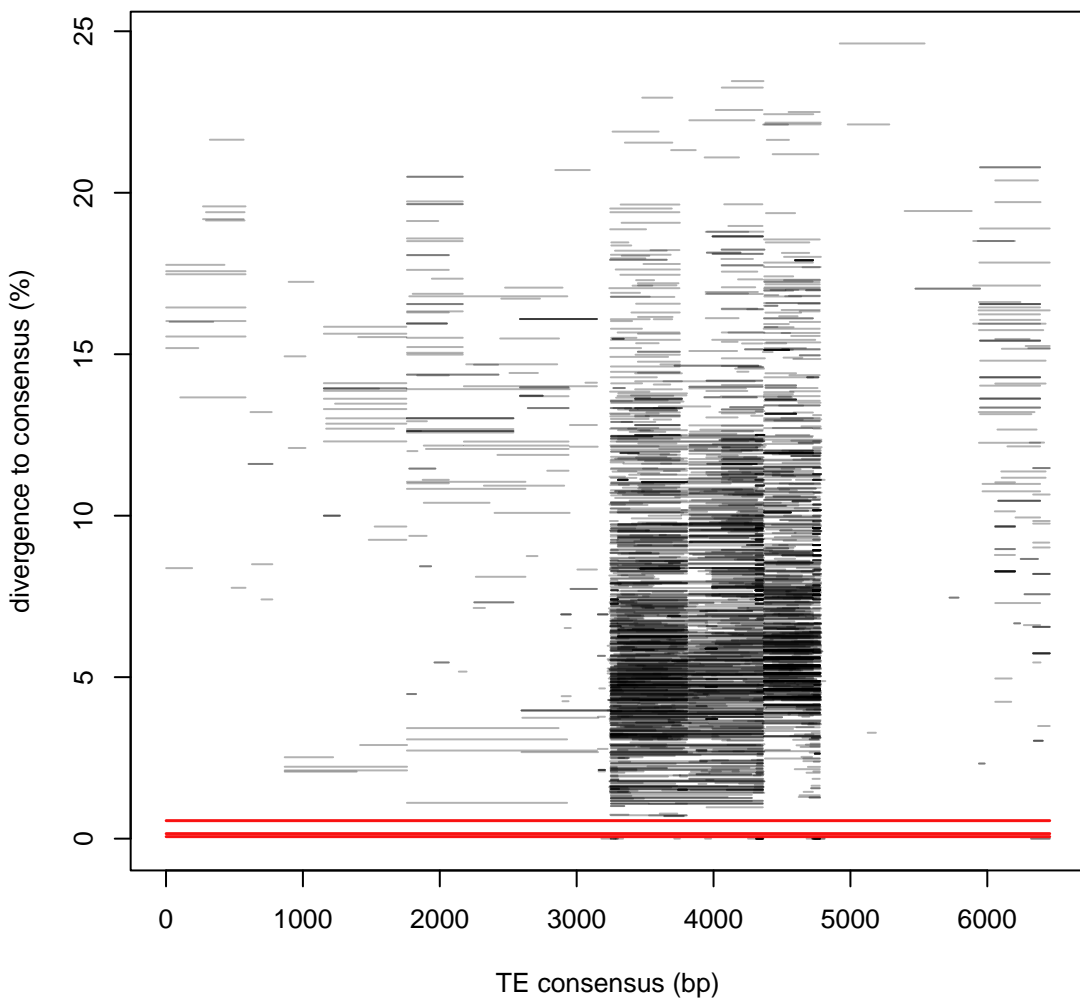
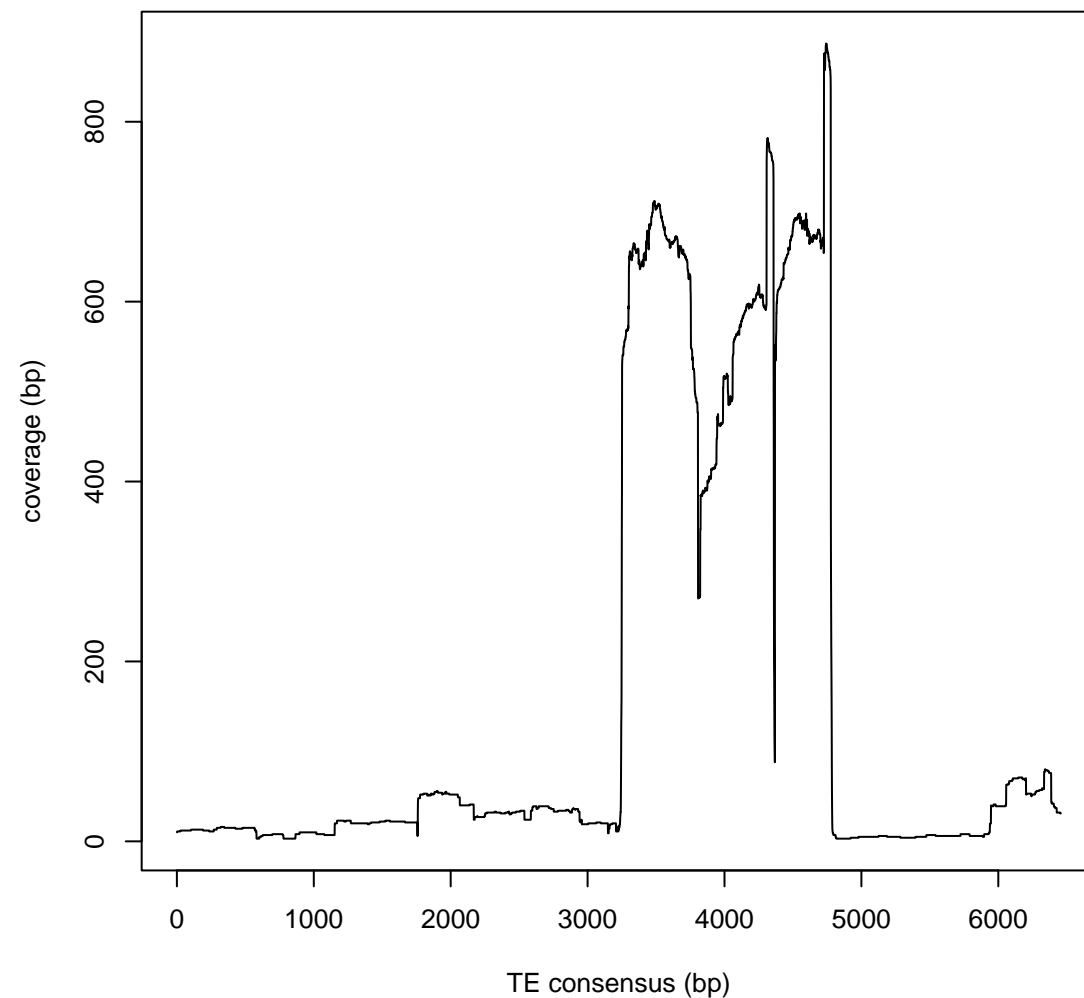


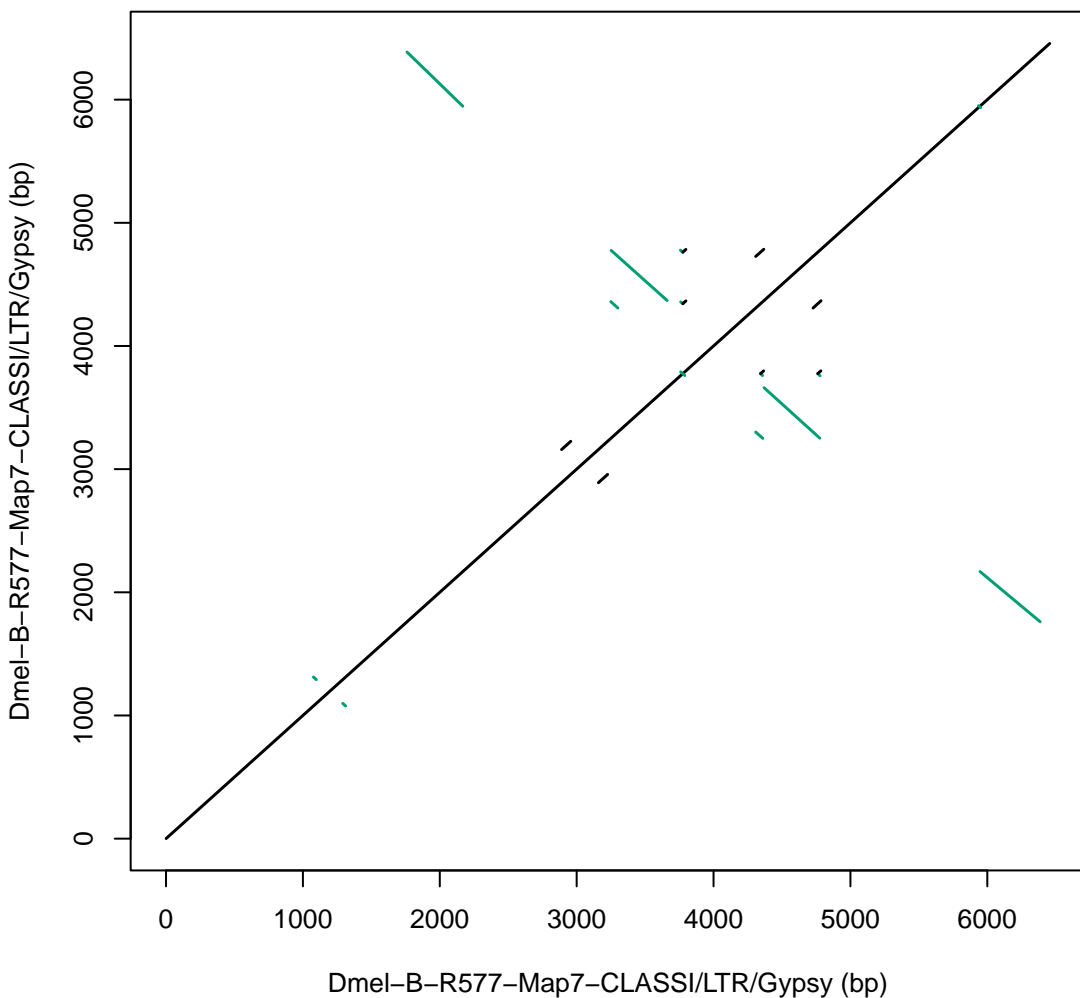
TE: Dmel-B-R577-Map7-CLASSI/LTR/Gypsy  
consensus size: 6456bp; fragments: 3294; full length: 3 (>=5810.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

