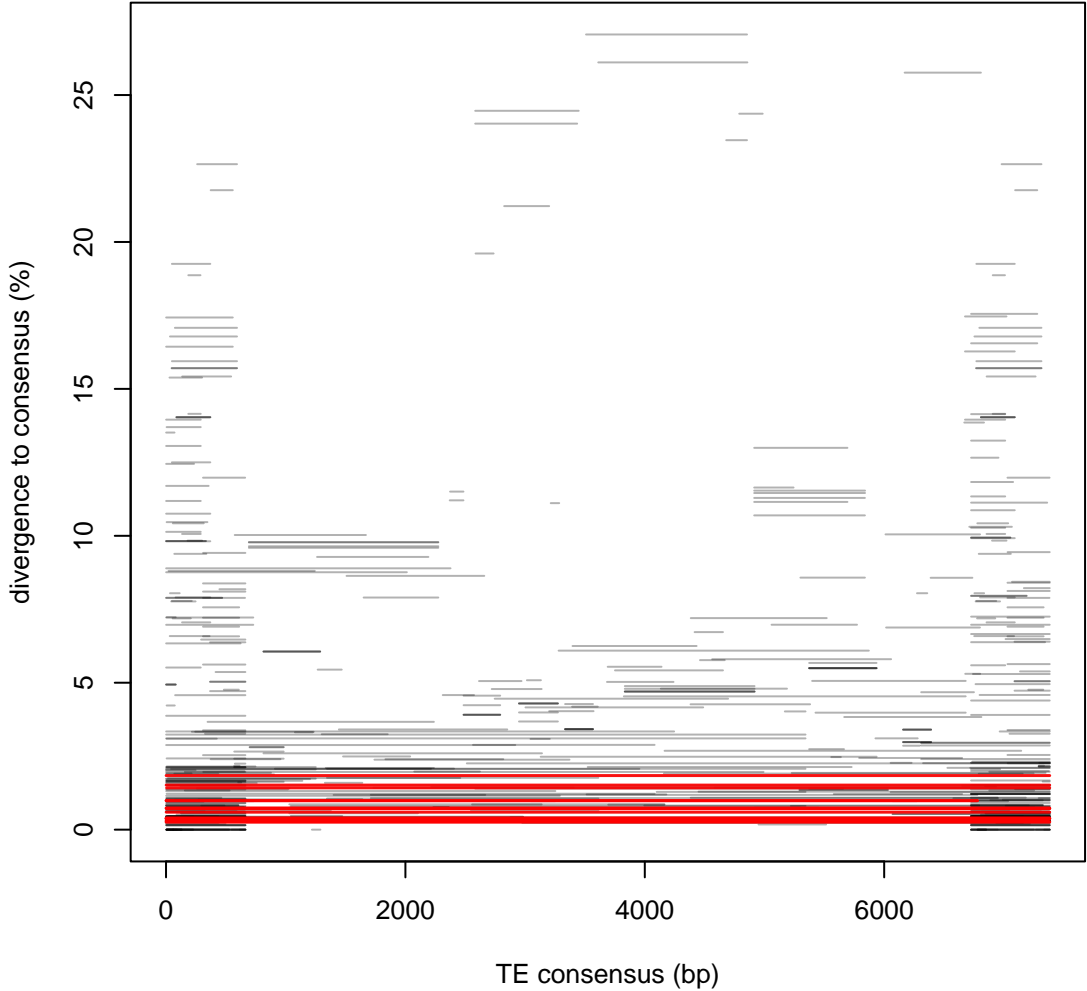
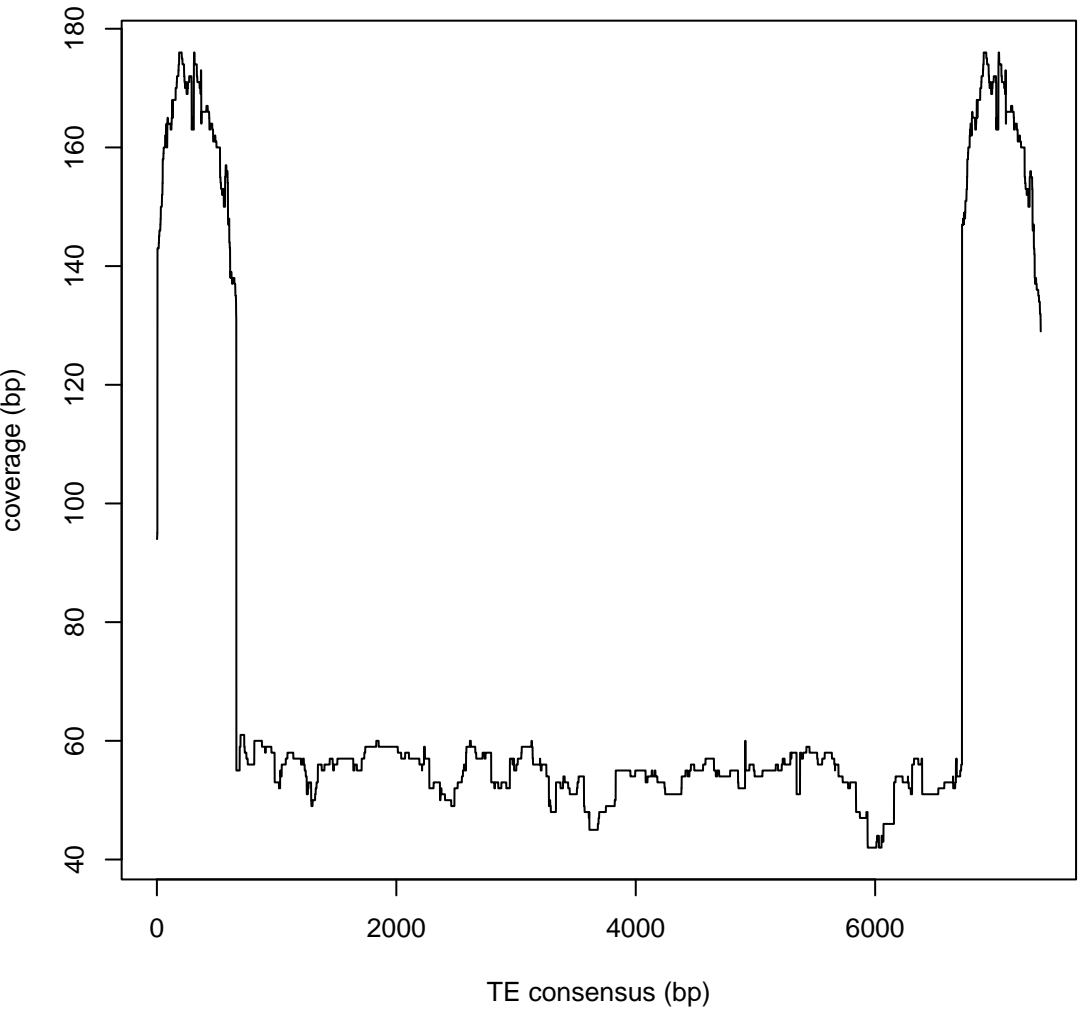


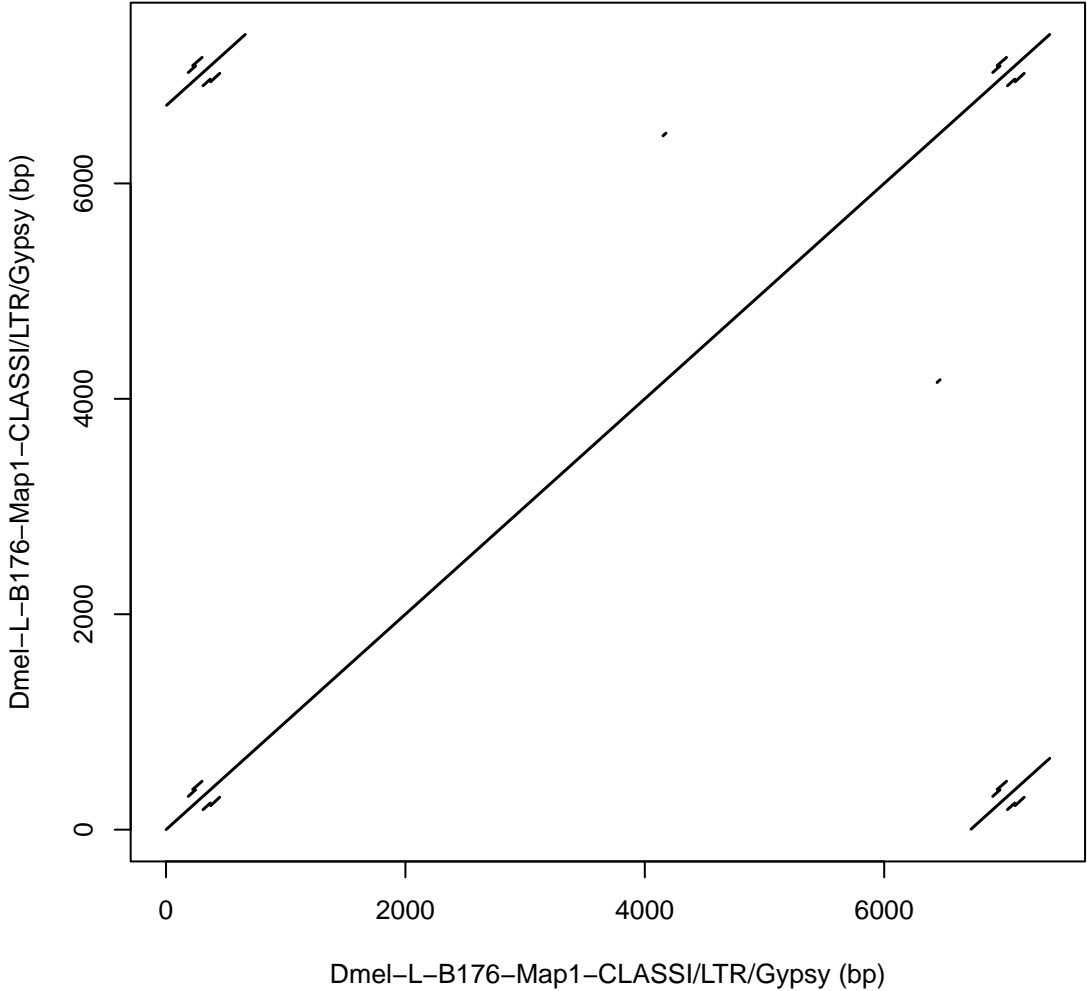
TE: Dmel-L-B176-Map1-CLASSI/LTR/Gypsy  
consensus size: 7383bp; fragments: 585; full length: 16 (>=6644.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

