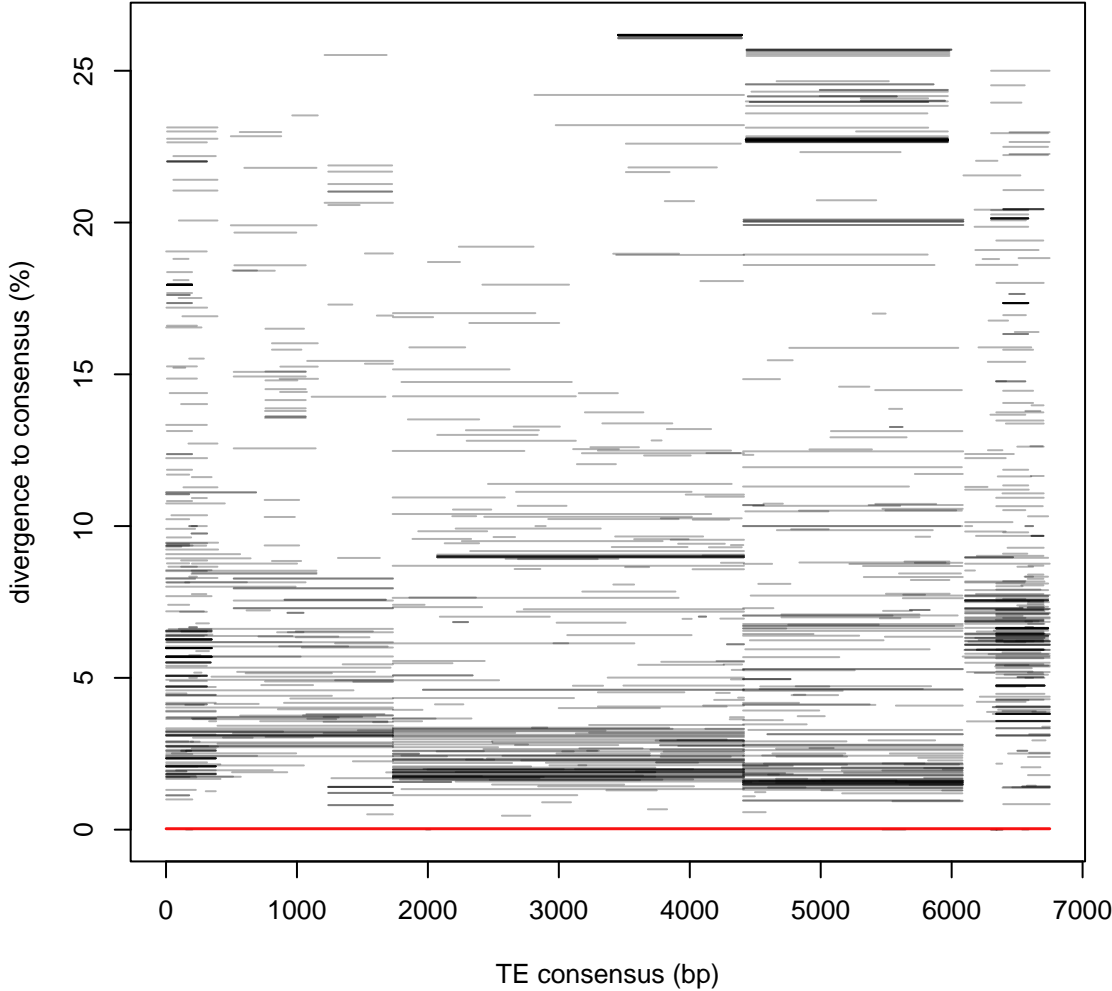
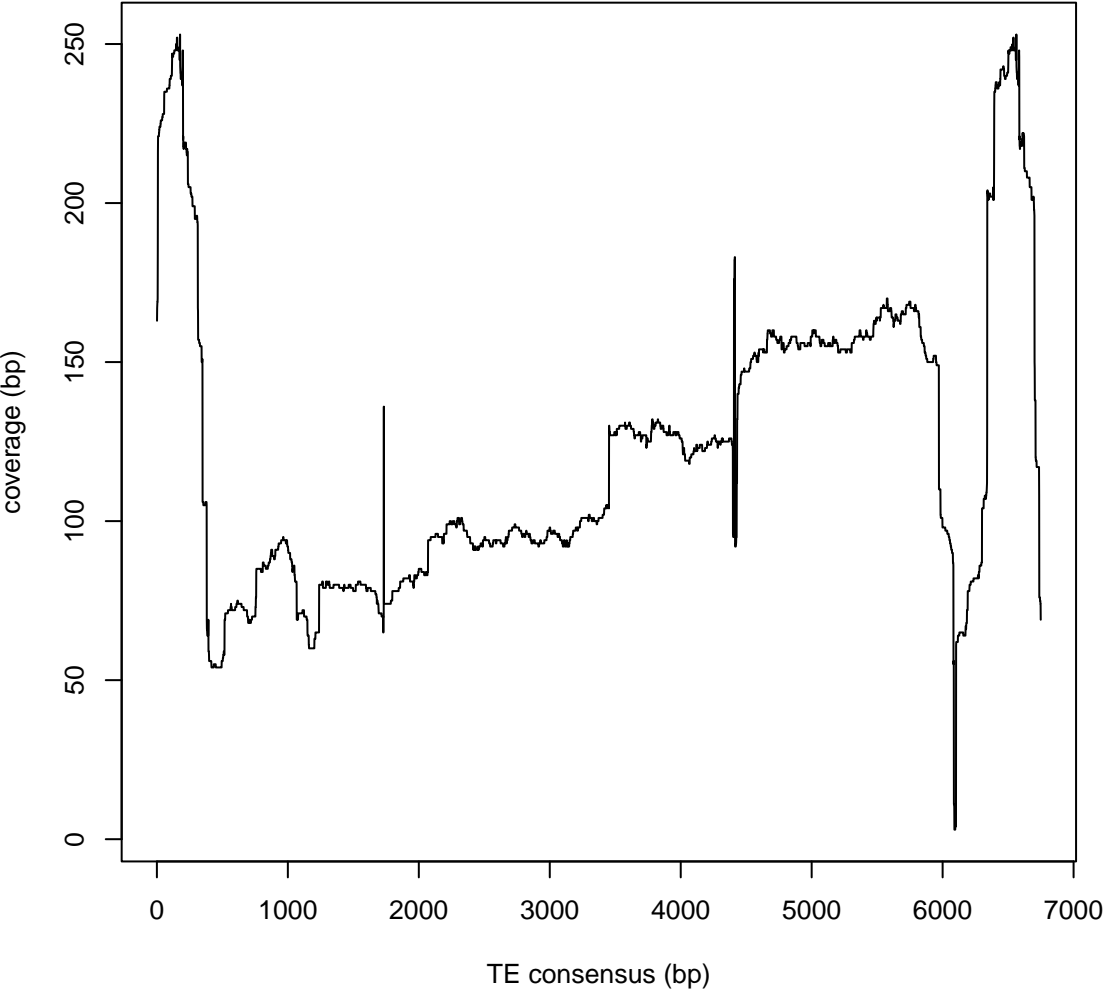


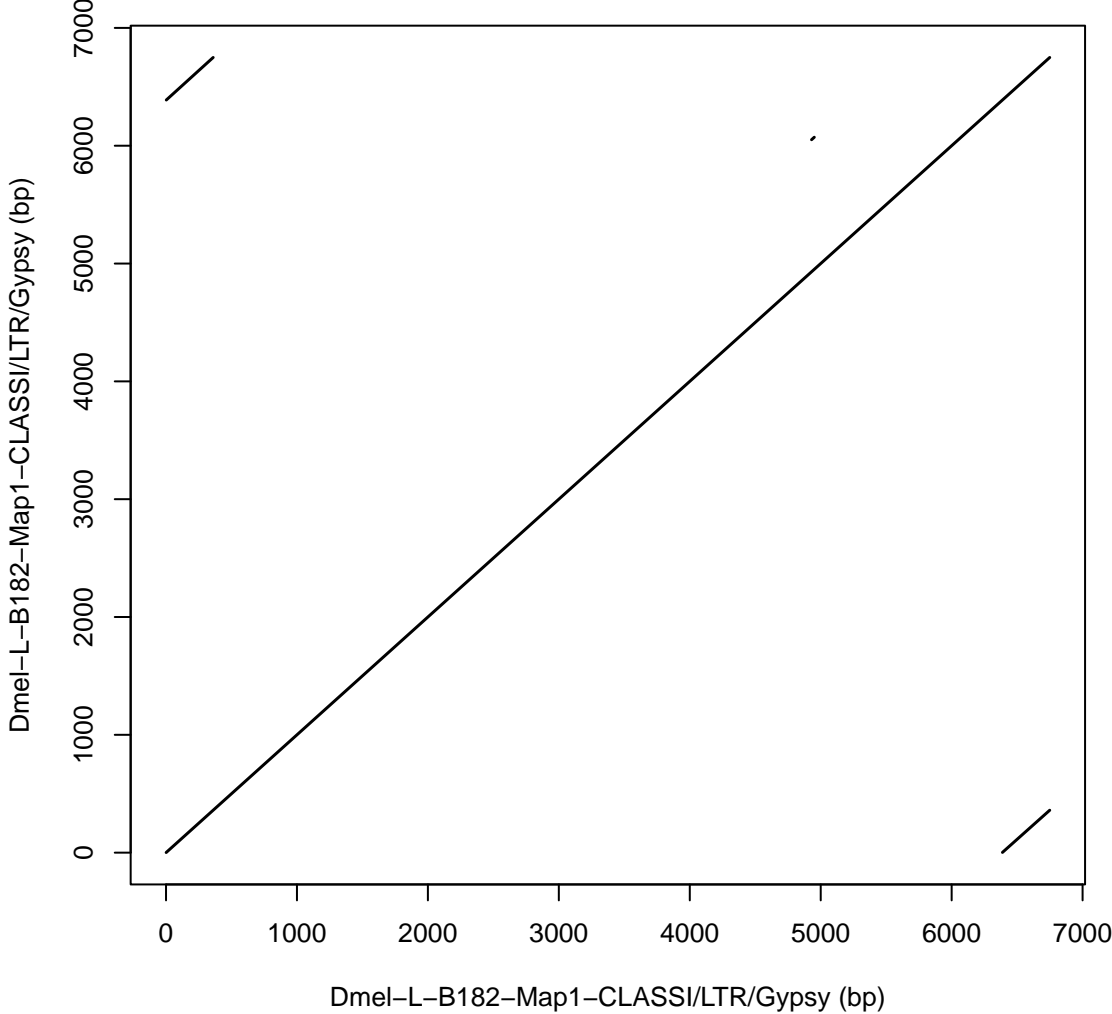
TE: Dmel-L-B182-Map1-CLASSI/LTR/Gypsy  
consensus size: 6749bp; fragments: 1206; full length: 1 (>=6074.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

