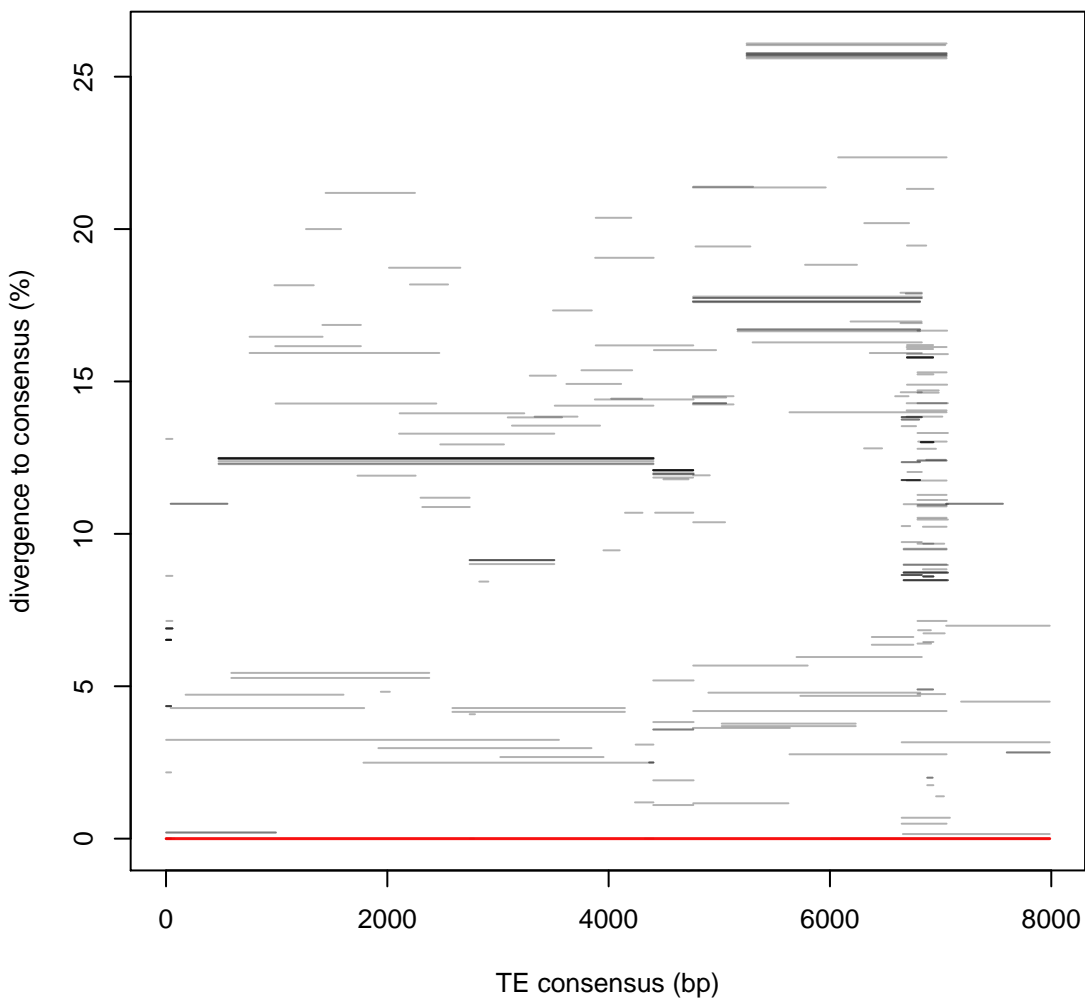
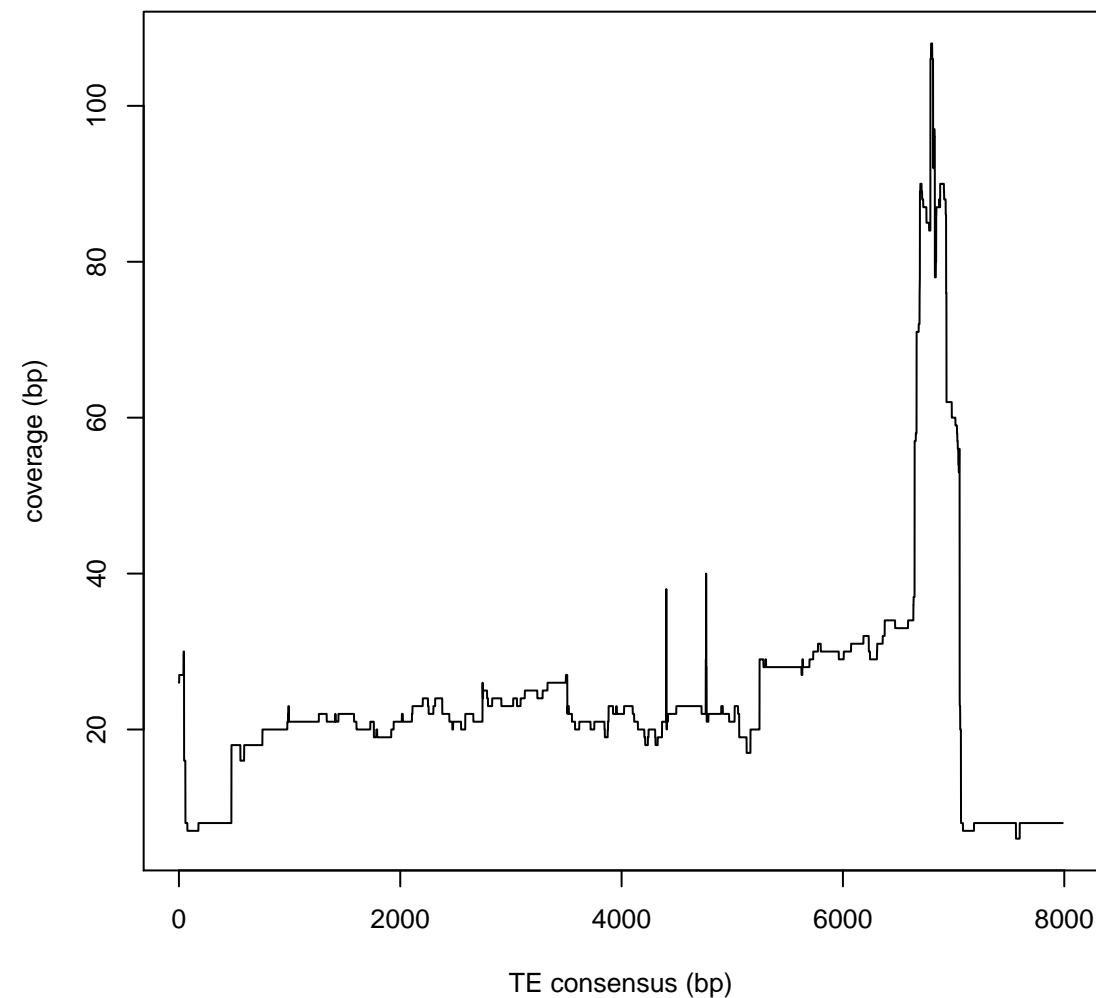


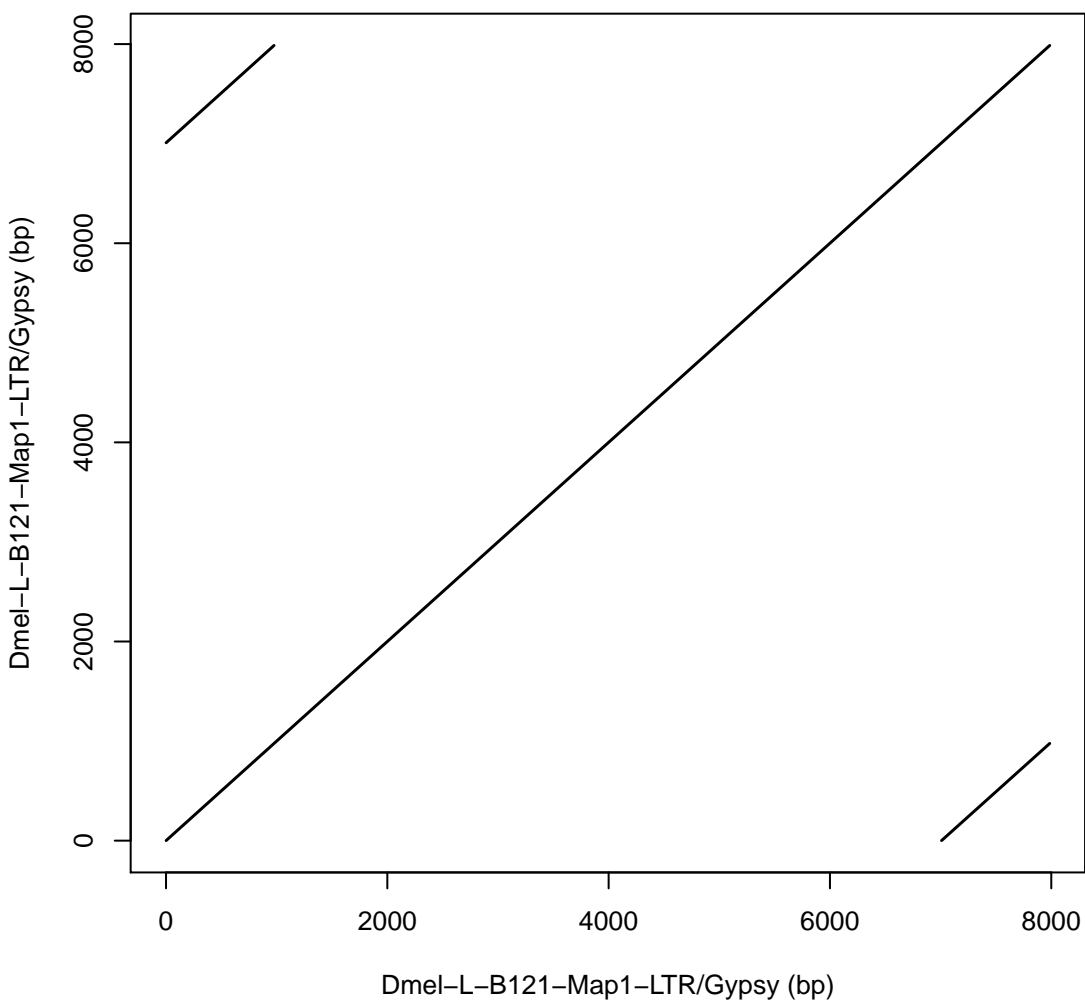
TE: Dmel-L-B121-Map1-LTR/Gypsy  
consensus size: 7986bp; fragments: 267; full length: 1 (>=7187.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

