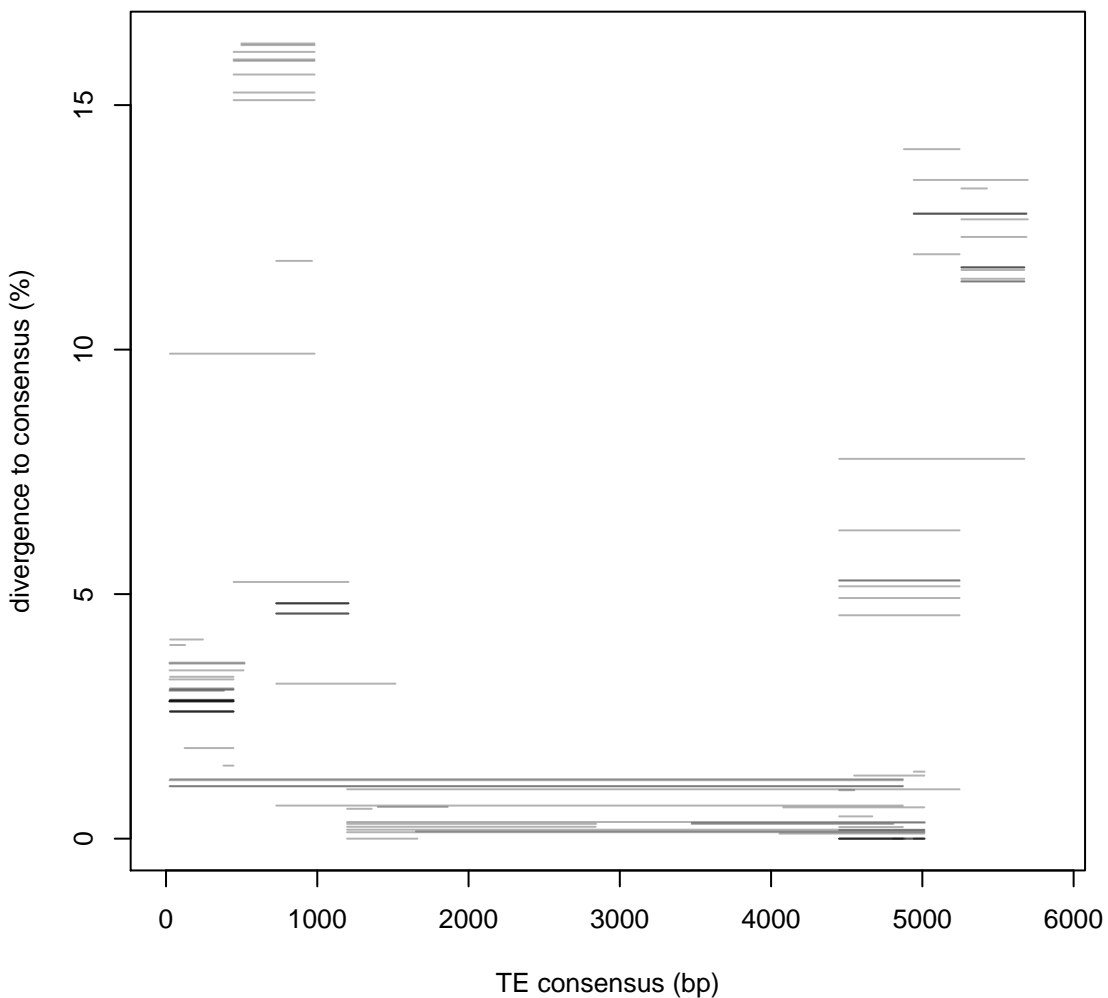
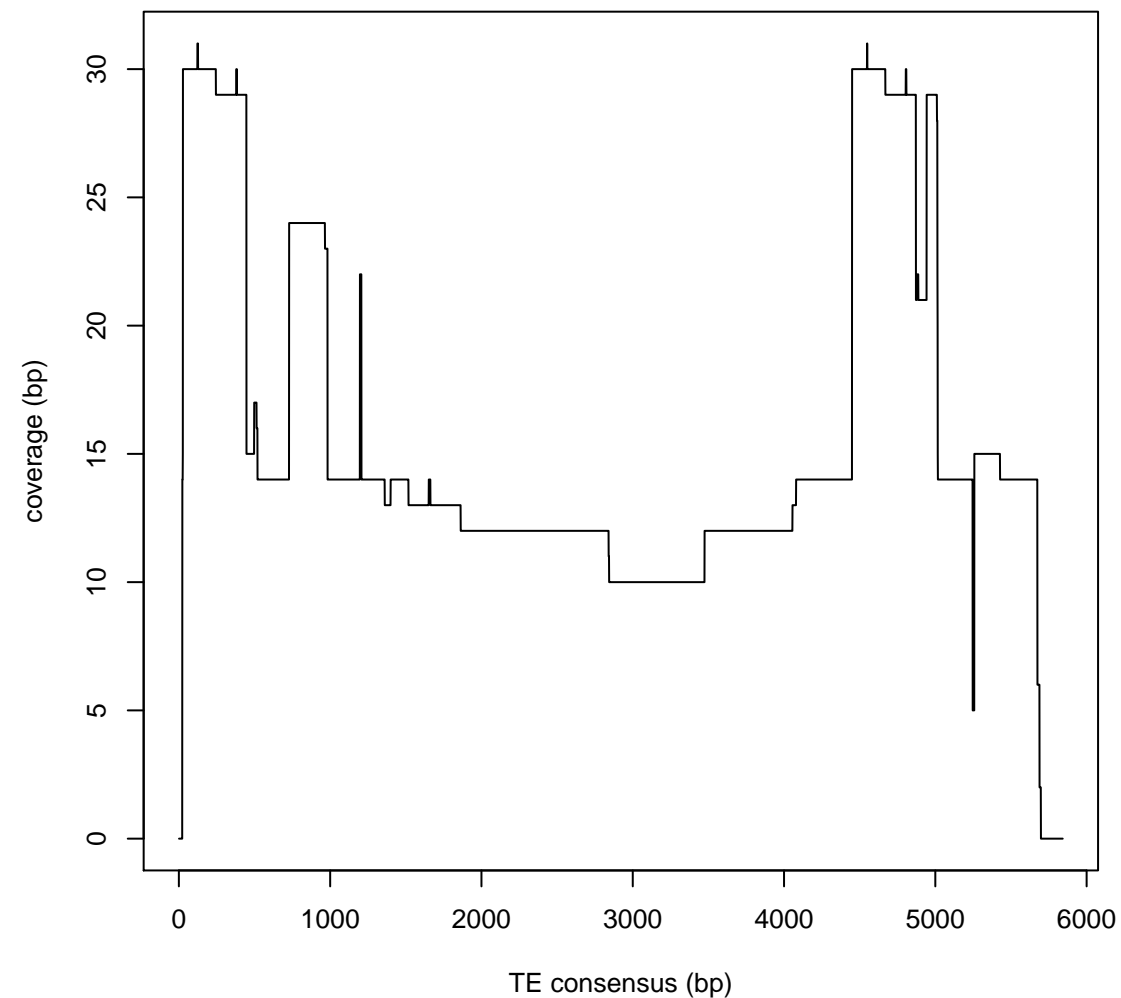


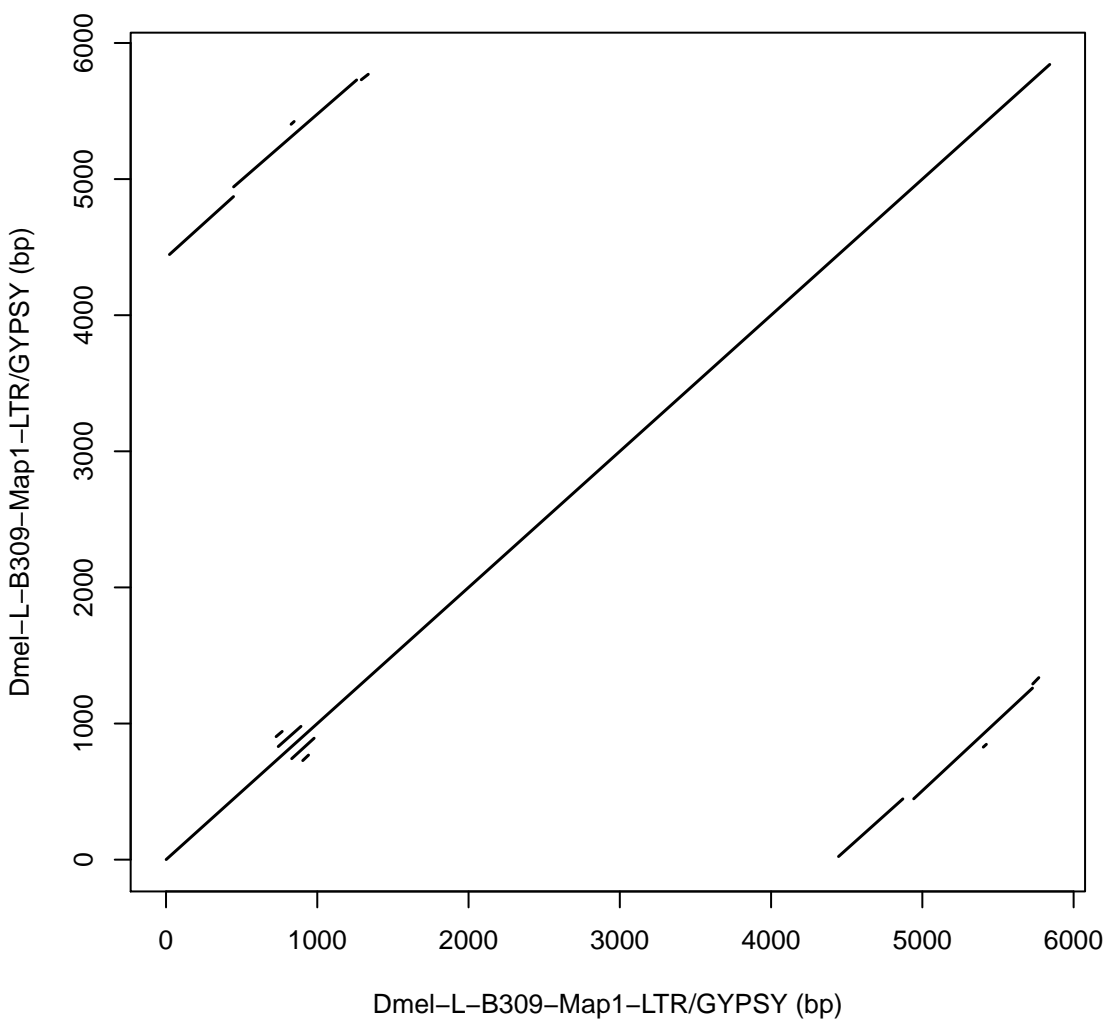
TE: Dmel-L-B309-Map1-LTR/GYPsy  
consensus size: 5842bp; fragments: 102; full length: 0 (>=5257.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

