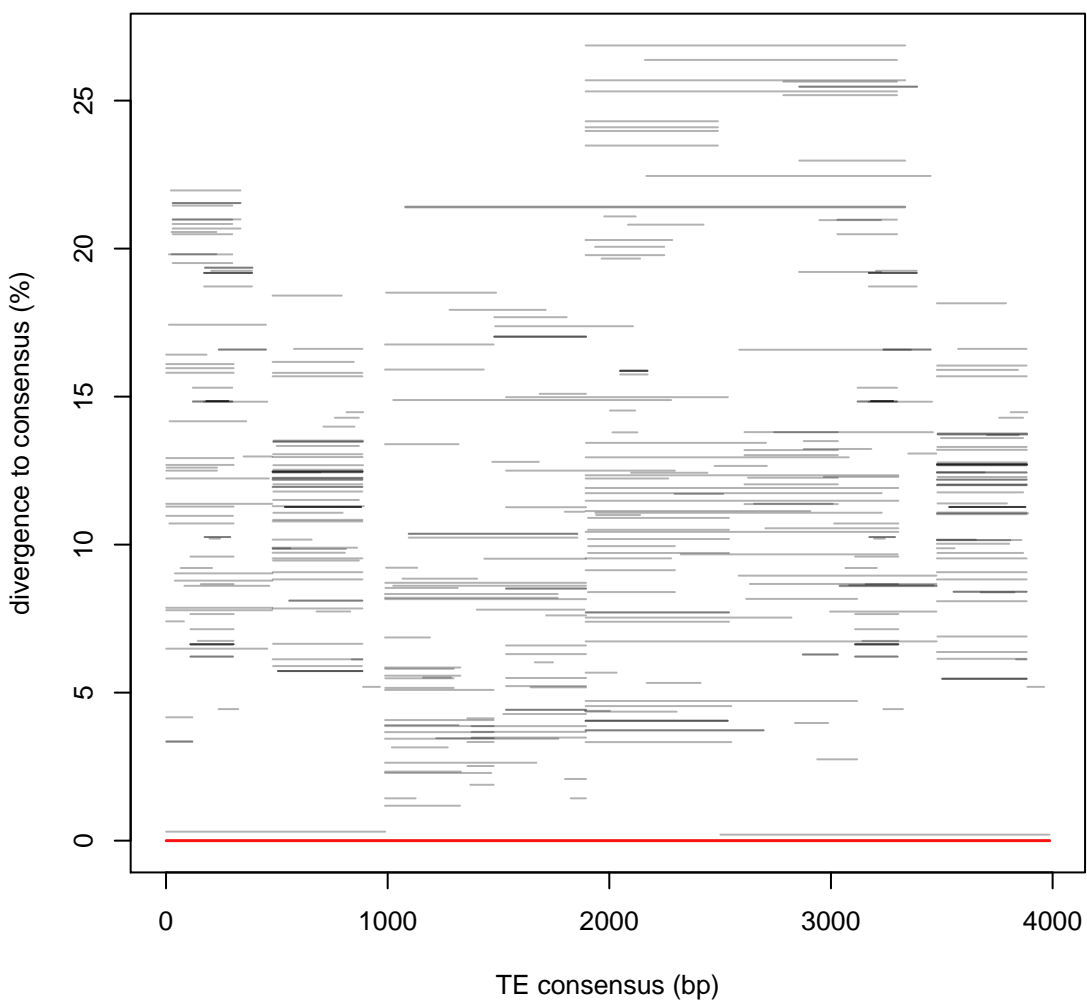
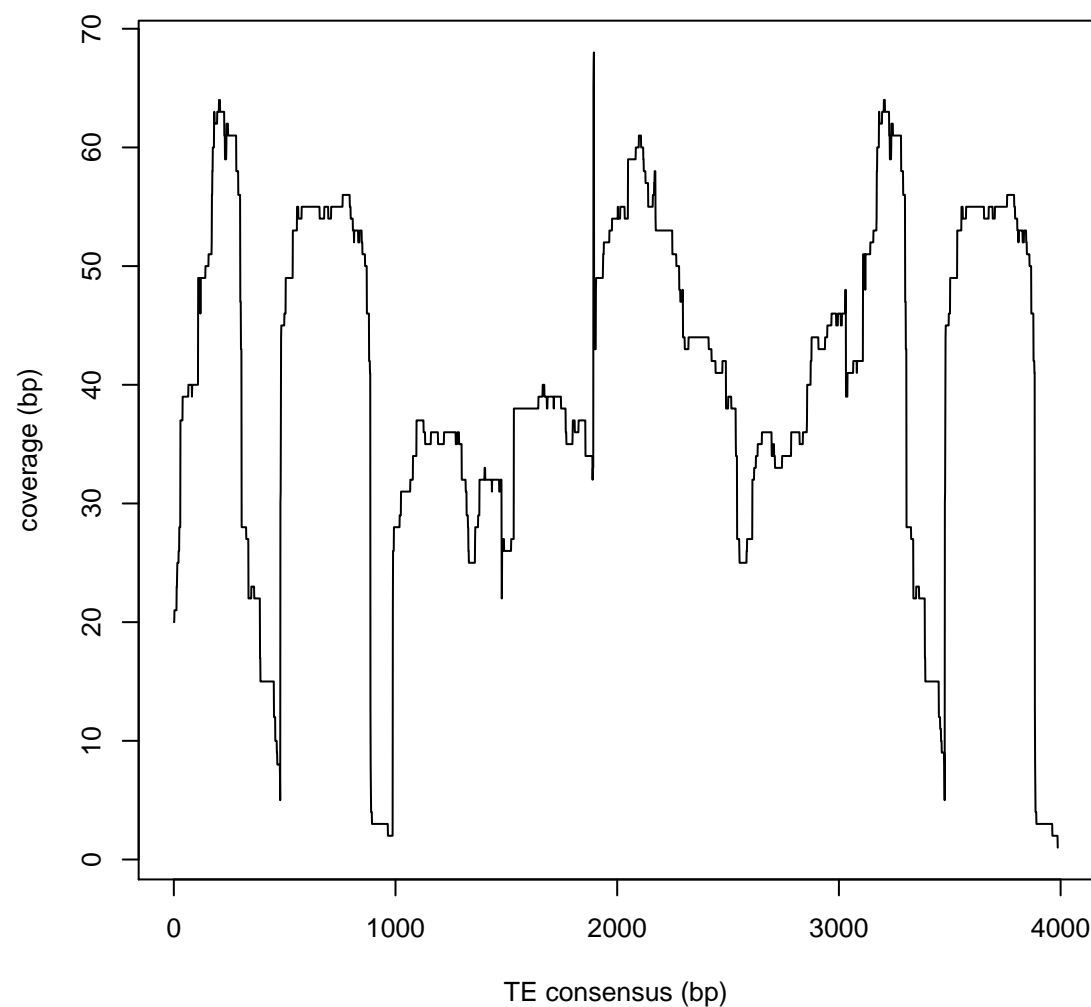


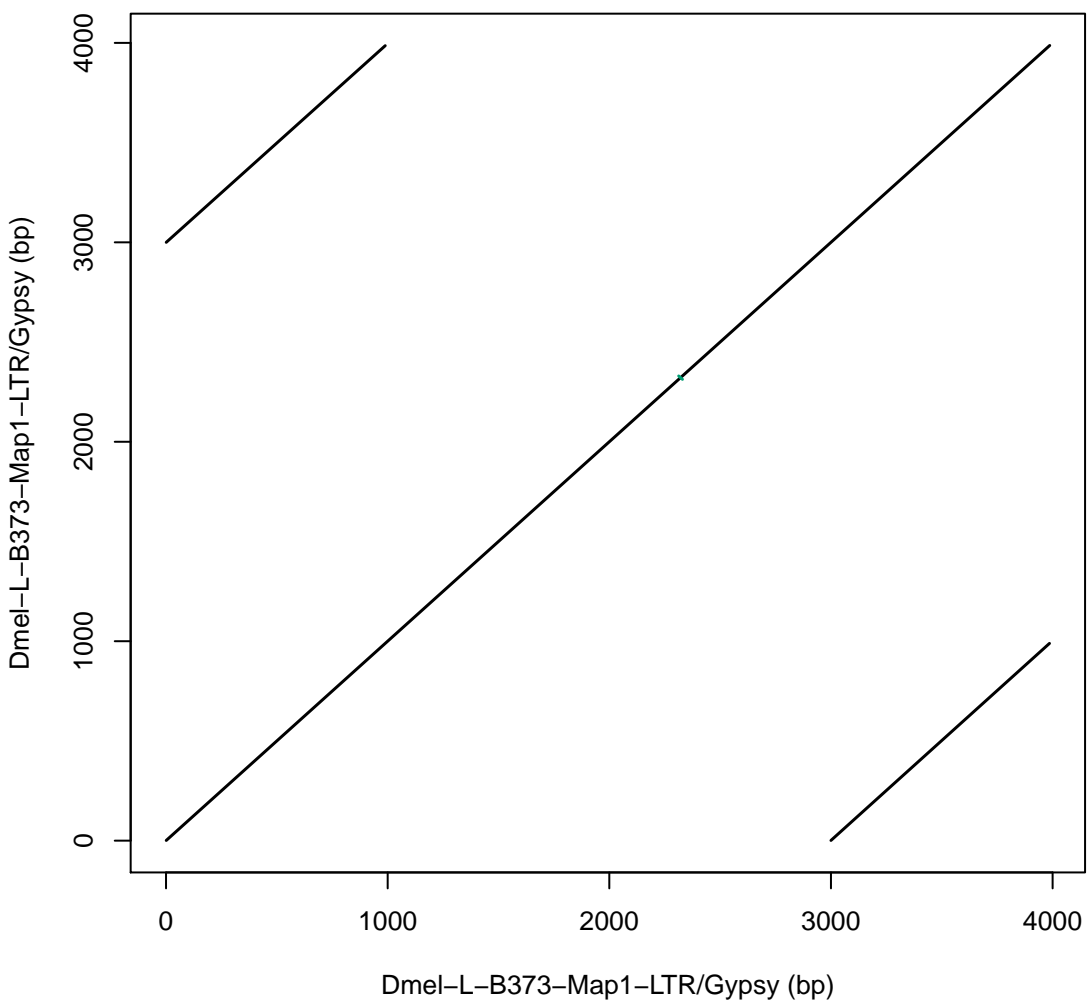
TE: Dmel-L-B373-Map1-LTR/Gypsy  
consensus size: 3987bp; fragments: 394; full length: 1 (>=3588.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

