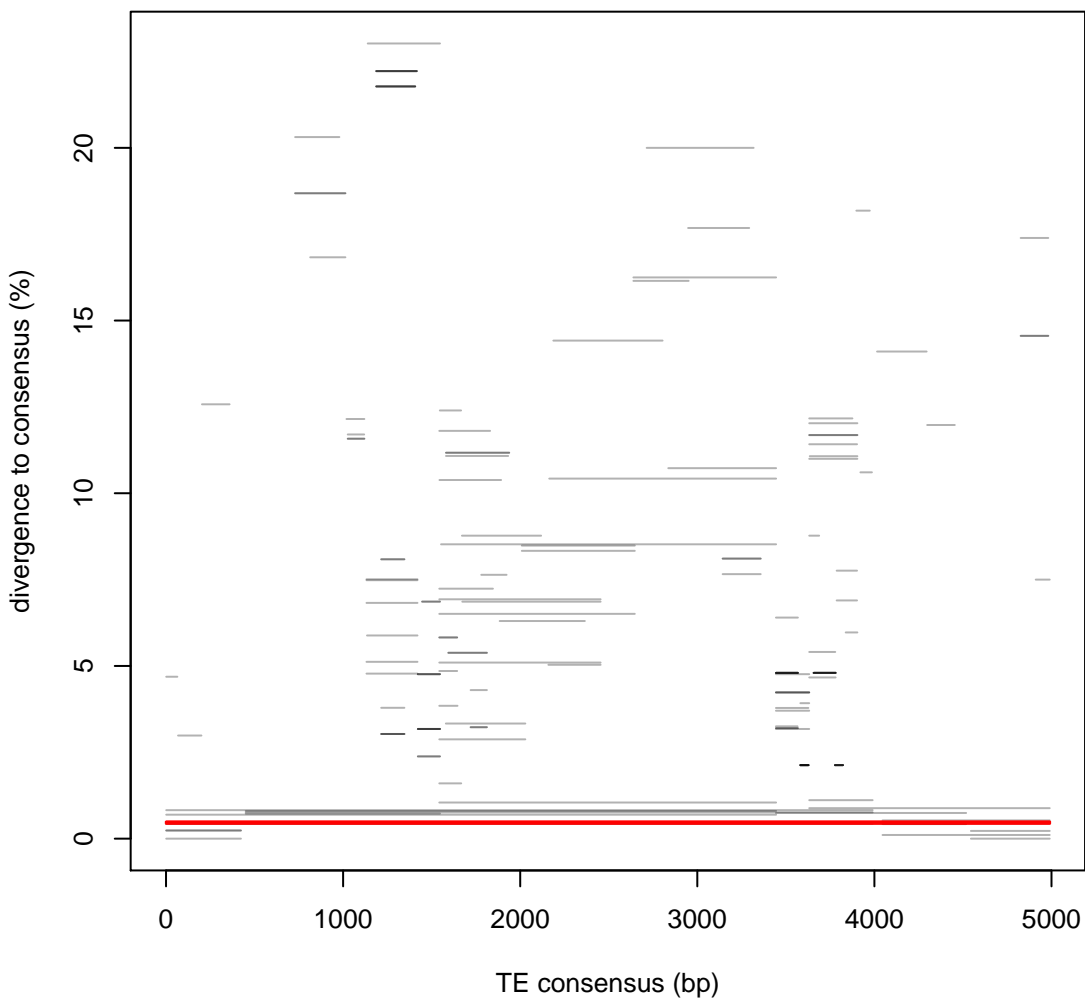
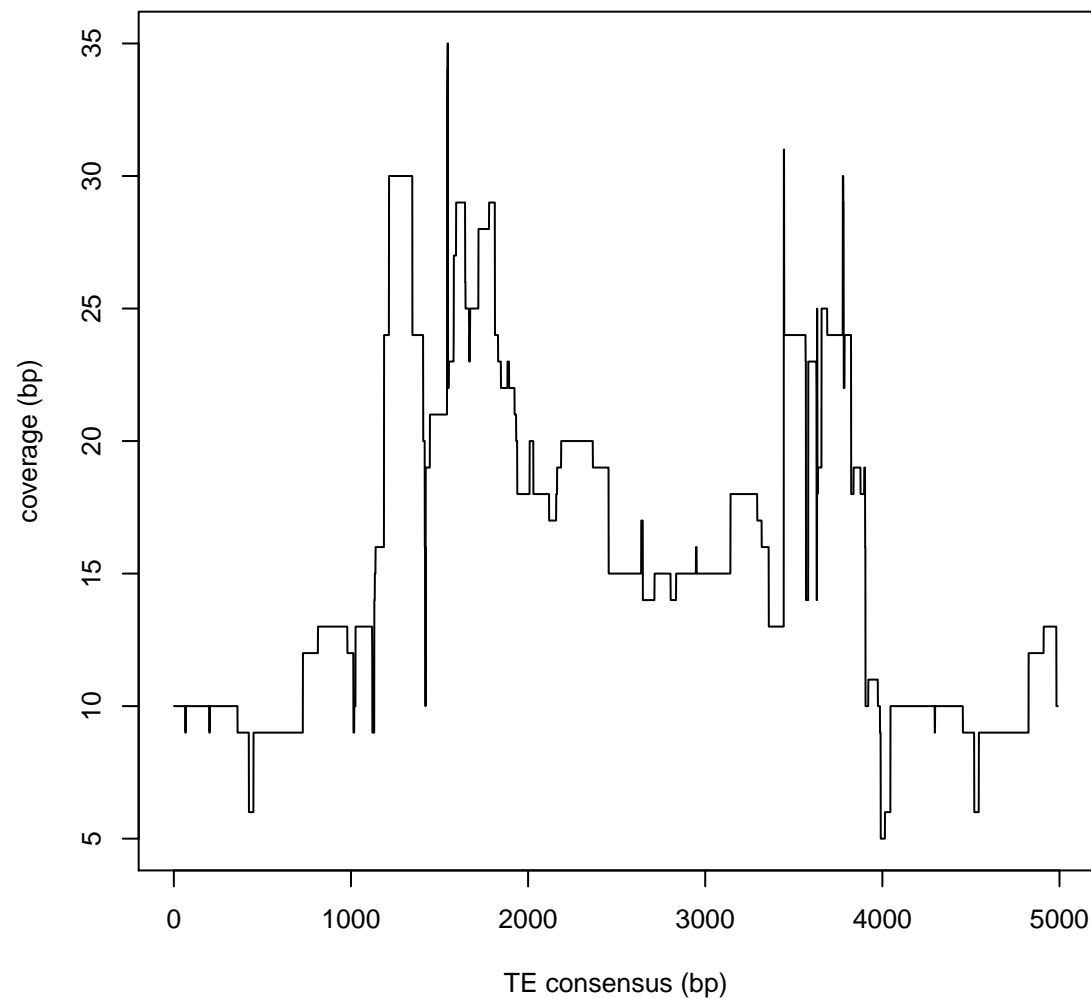


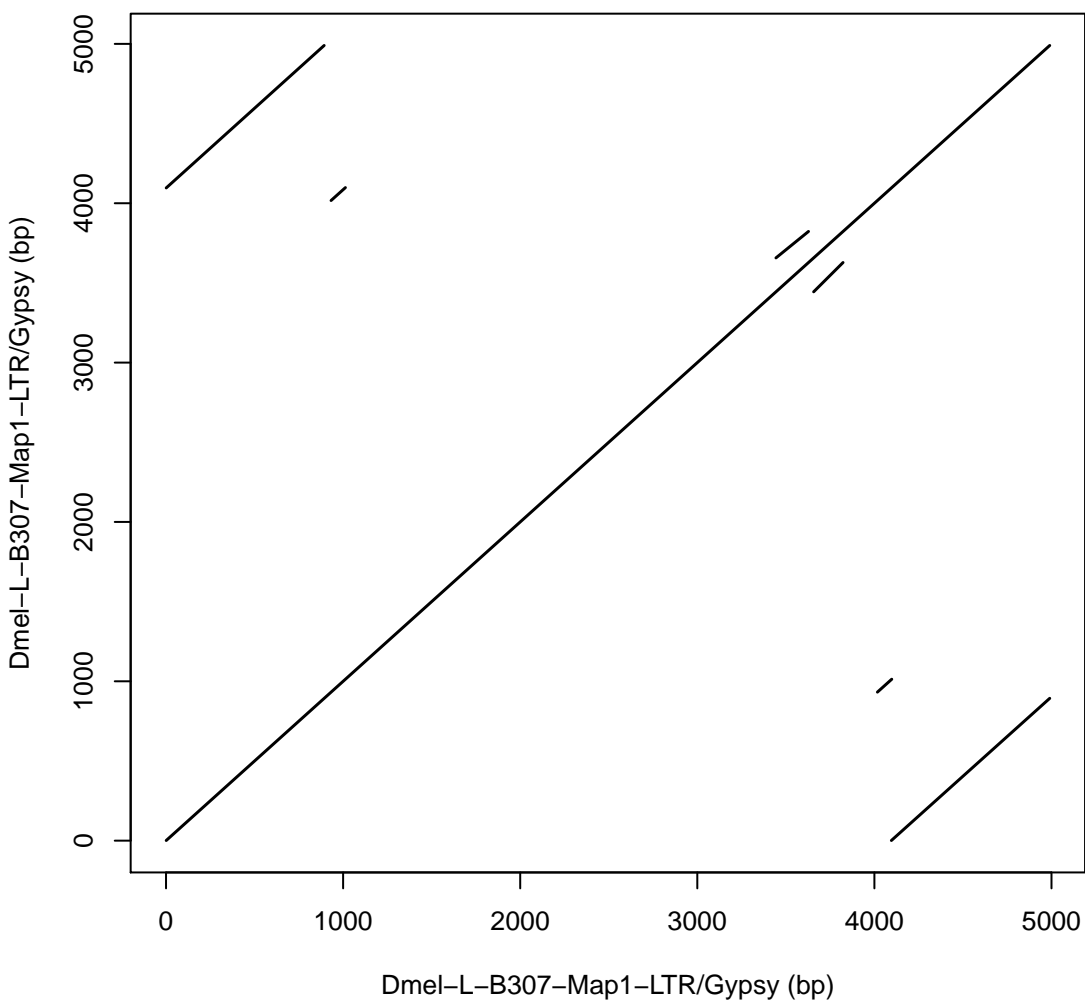
TE: Dmel-L-B307-Map1-LTR/Gypsy  
consensus size: 4990bp; fragments: 165; full length: 3 (>=4491bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

