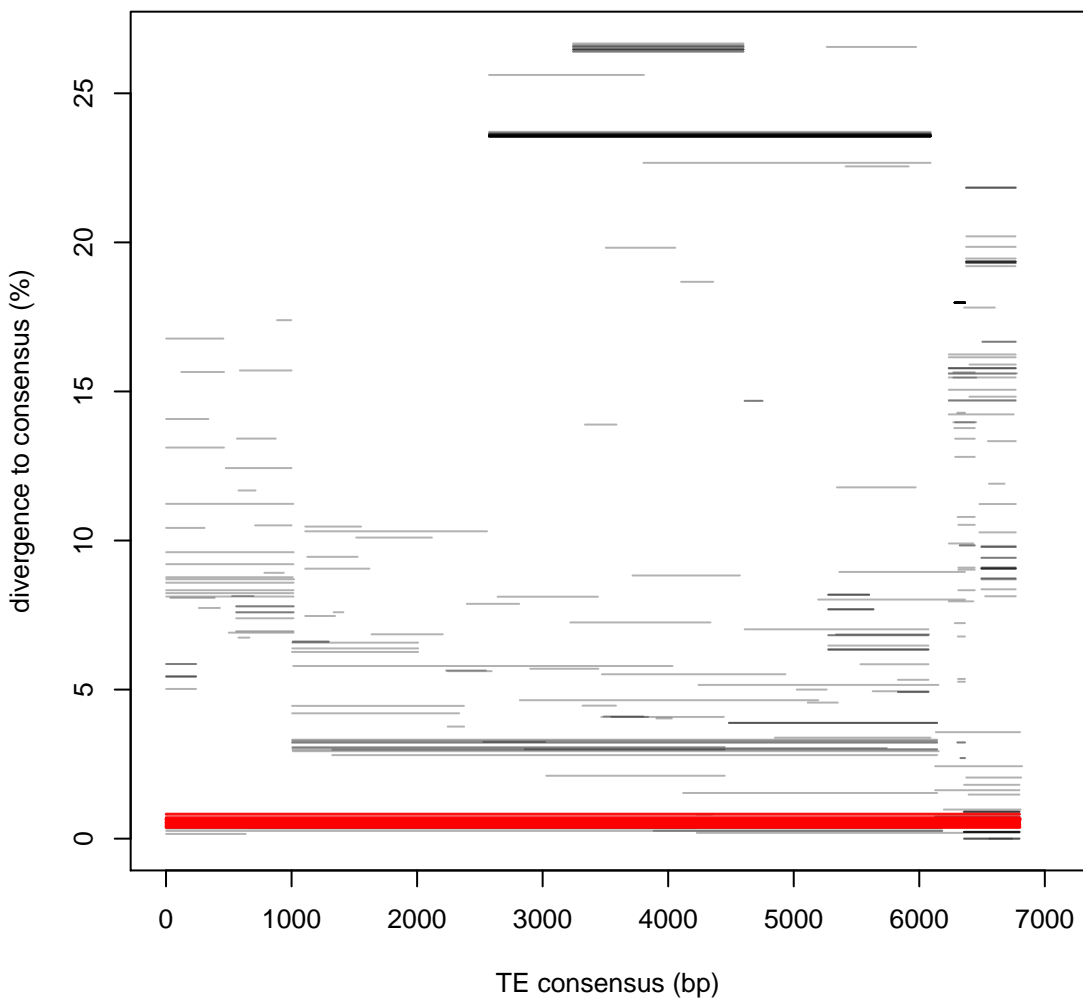
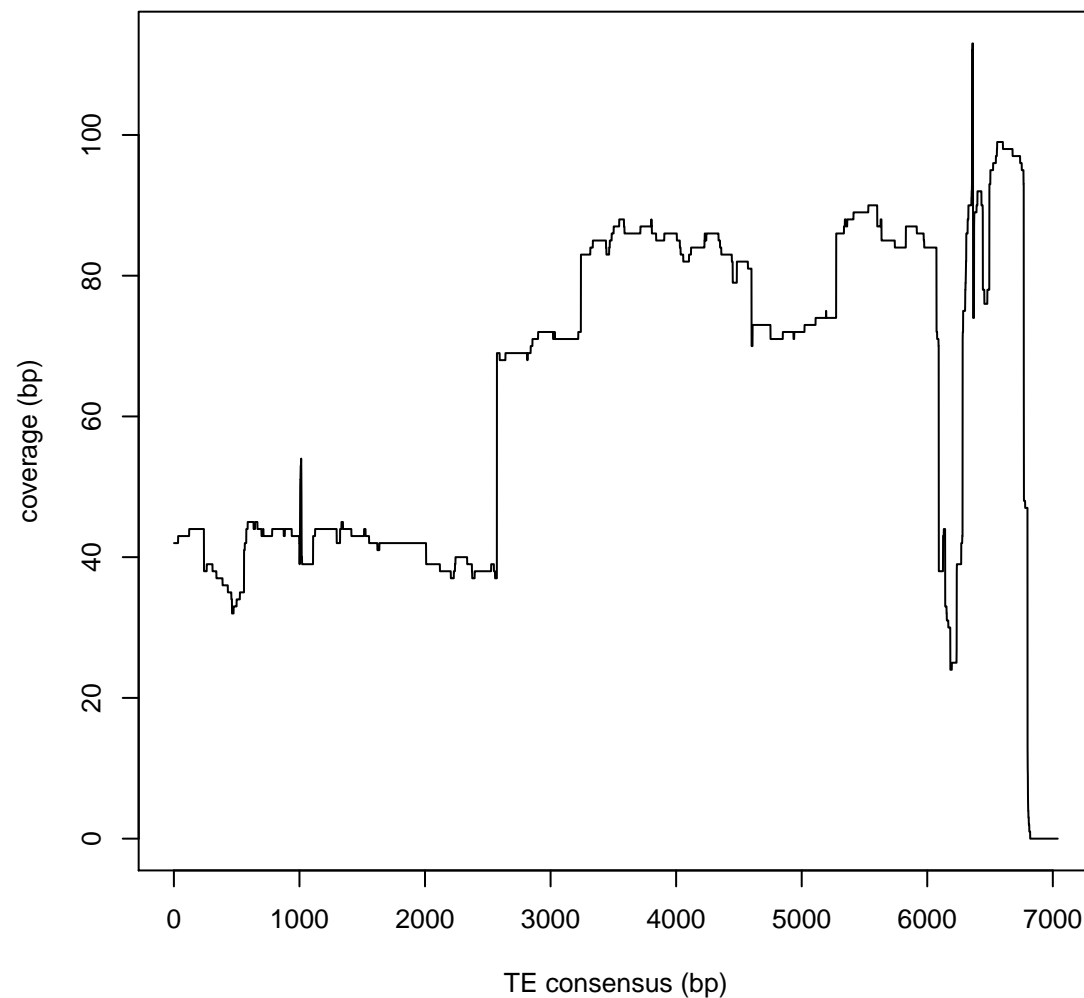


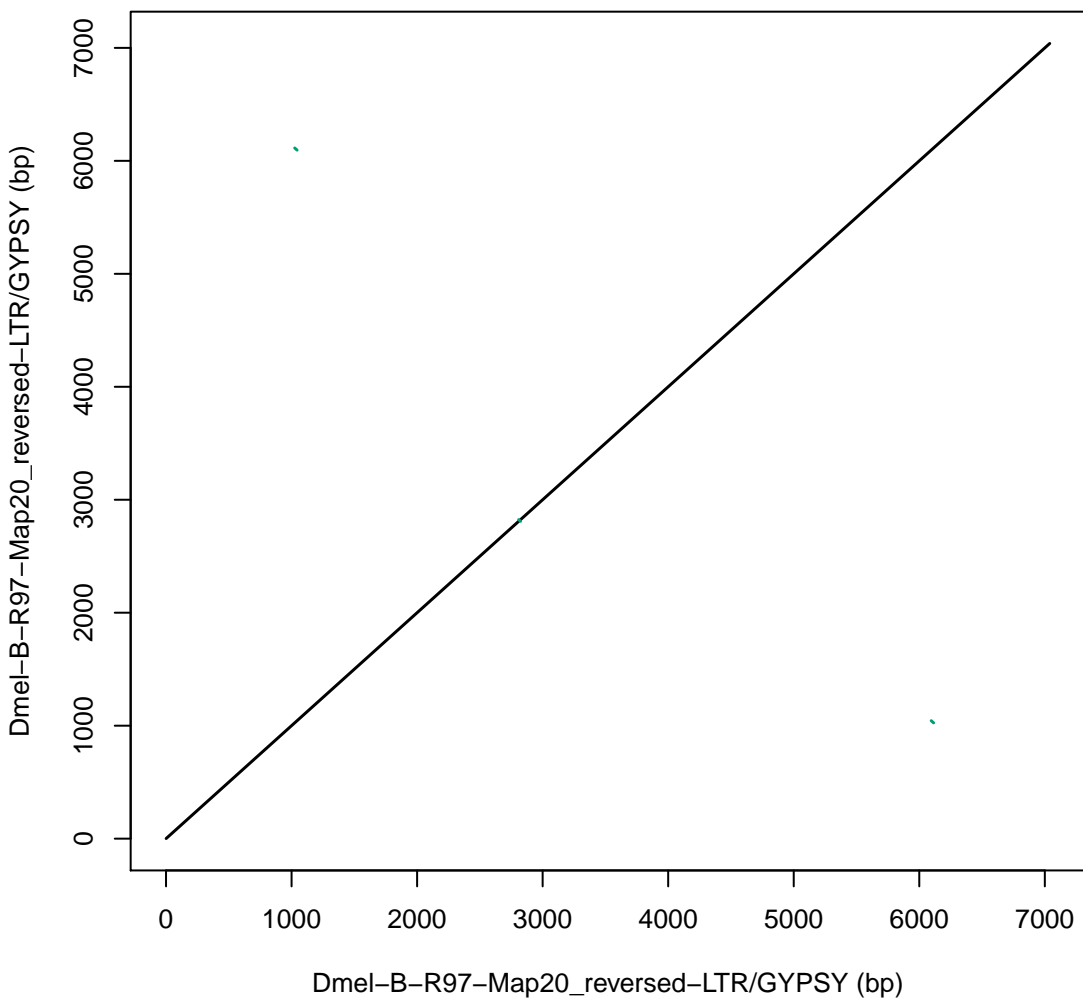
TE: Dmel-B-R97-Map20\_reversed-LTR/GYPSY  
consensus size: 7039bp; fragments: 328; full length: 15 (>=6335.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

