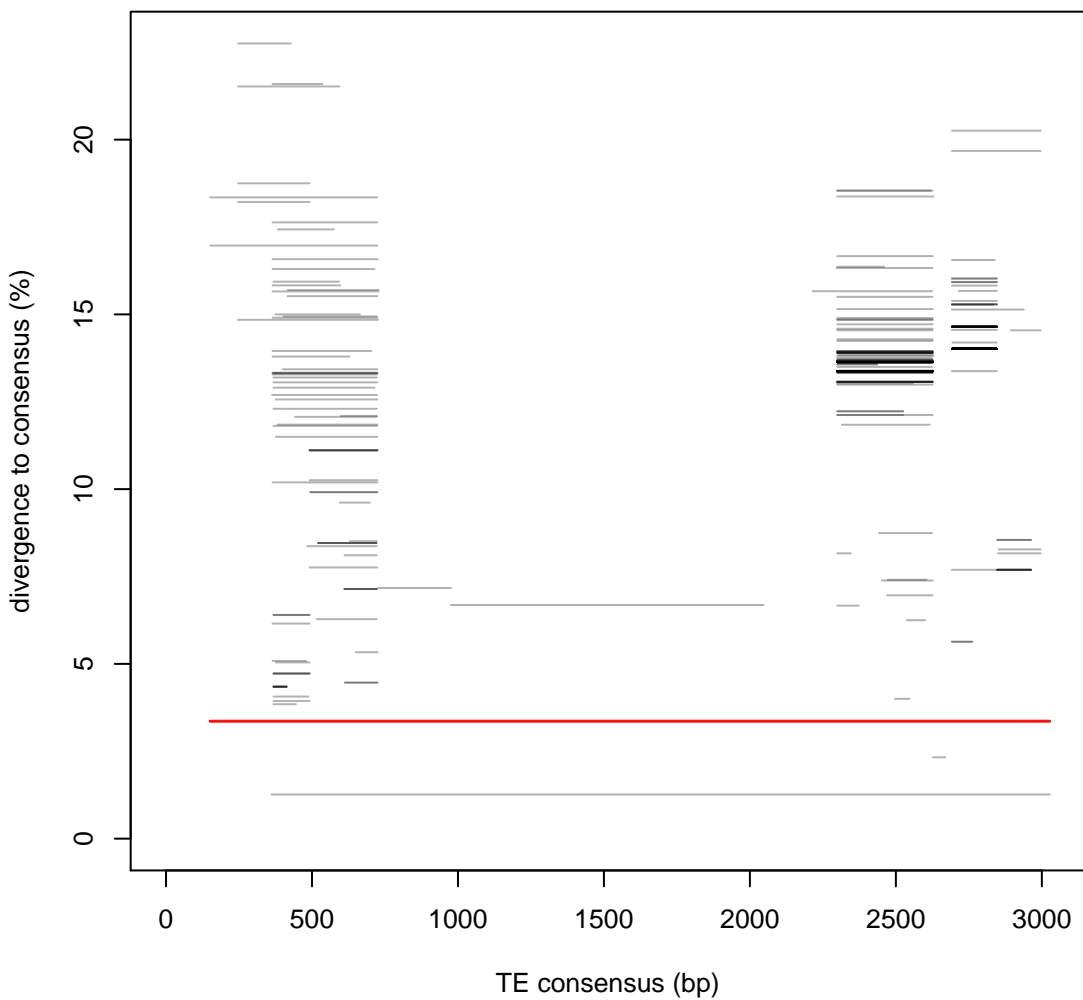
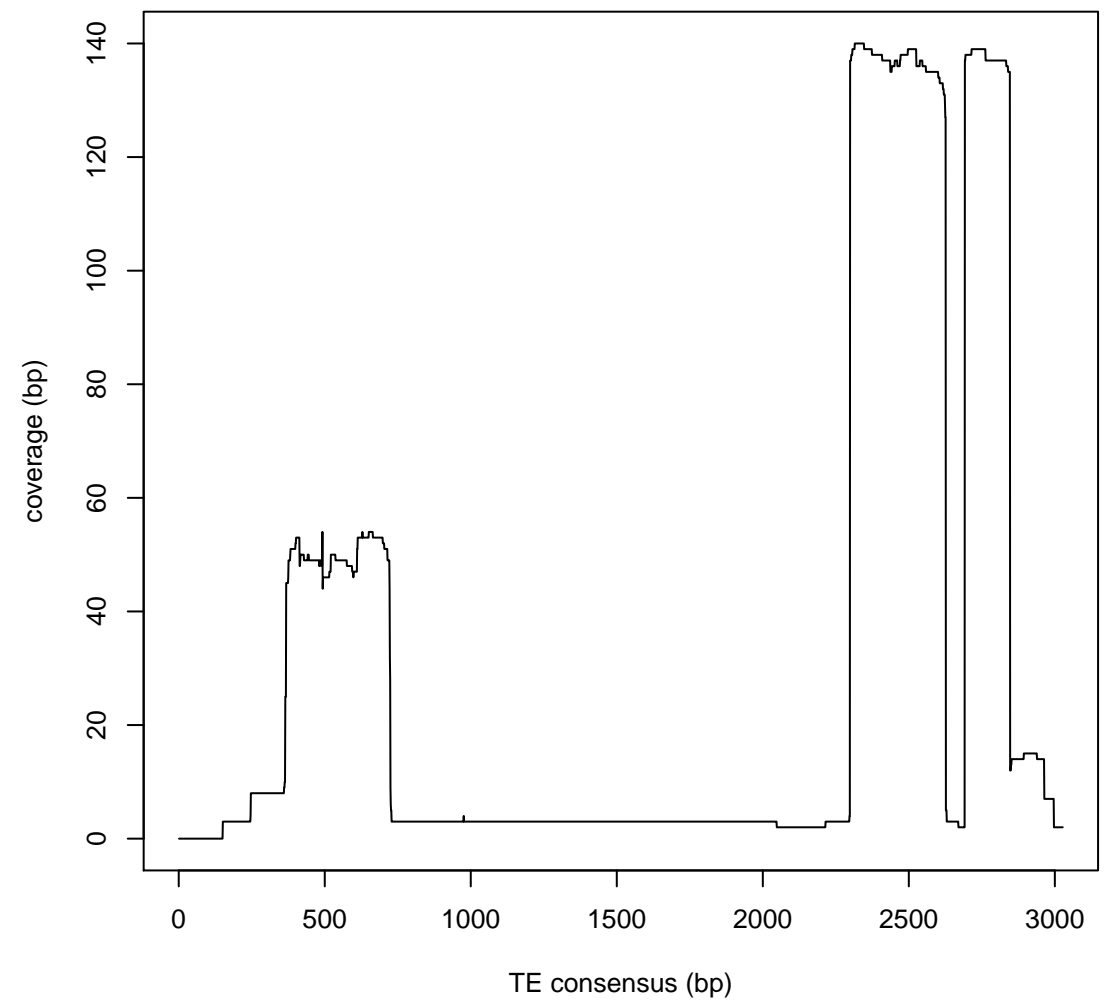


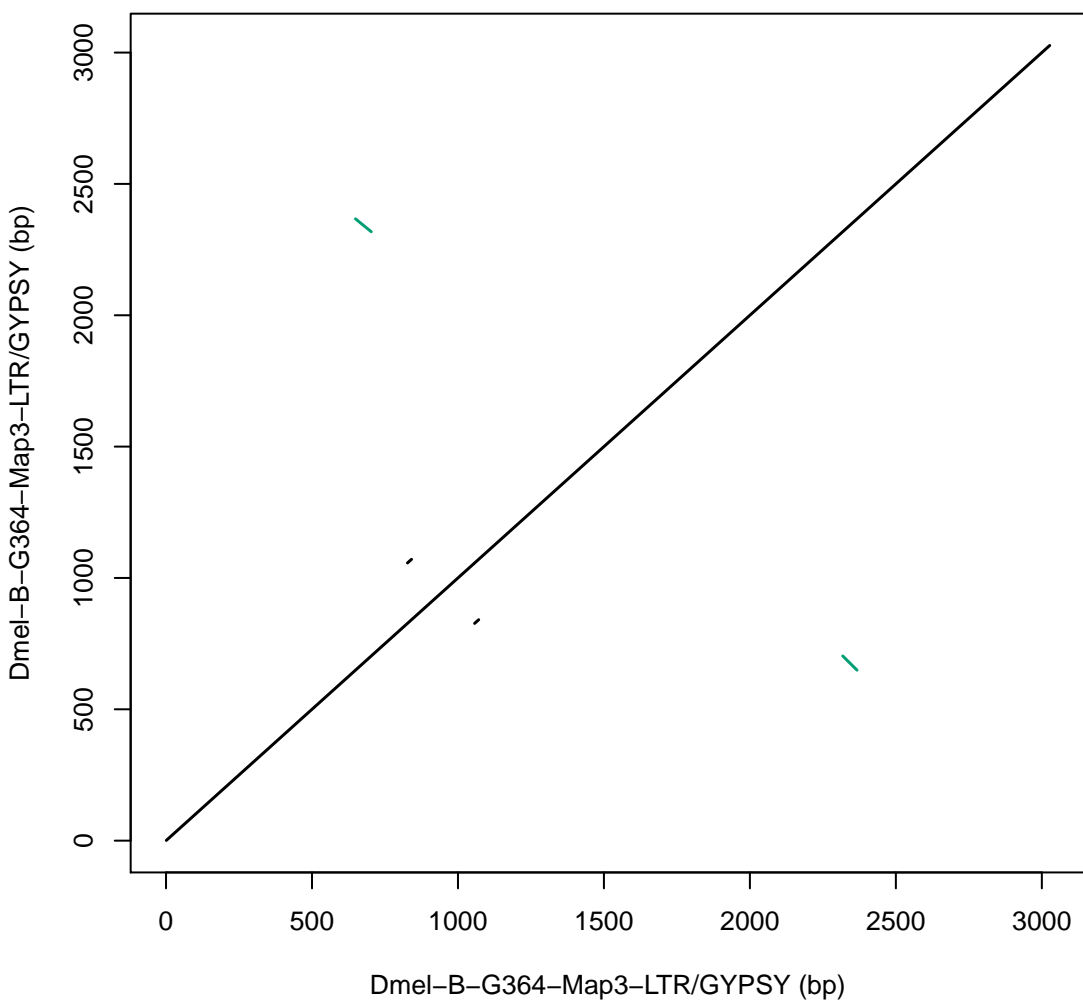
TE: Dmel-B-G364-Map3-LTR/GYPSY  
consensus size: 3027bp; fragments: 373; full length: 1 (>=2724.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

