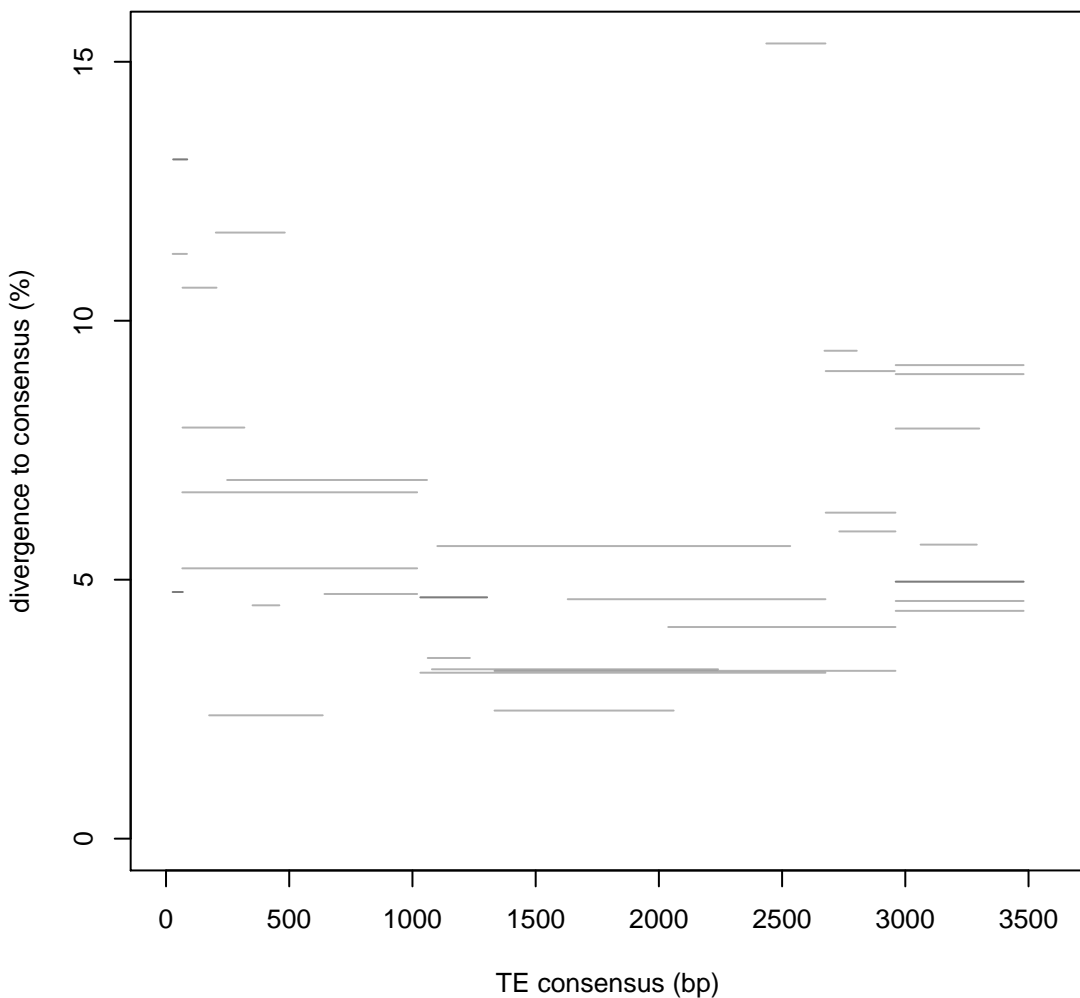
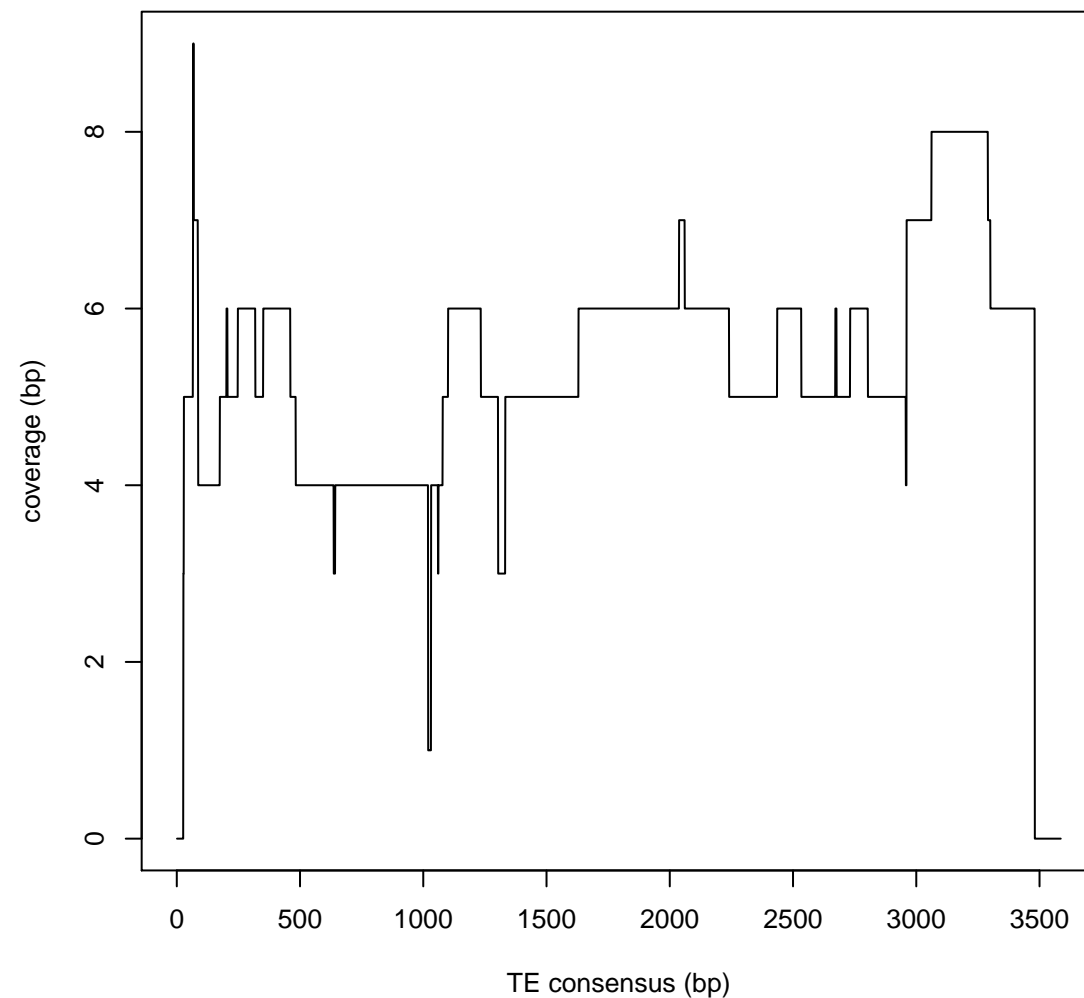


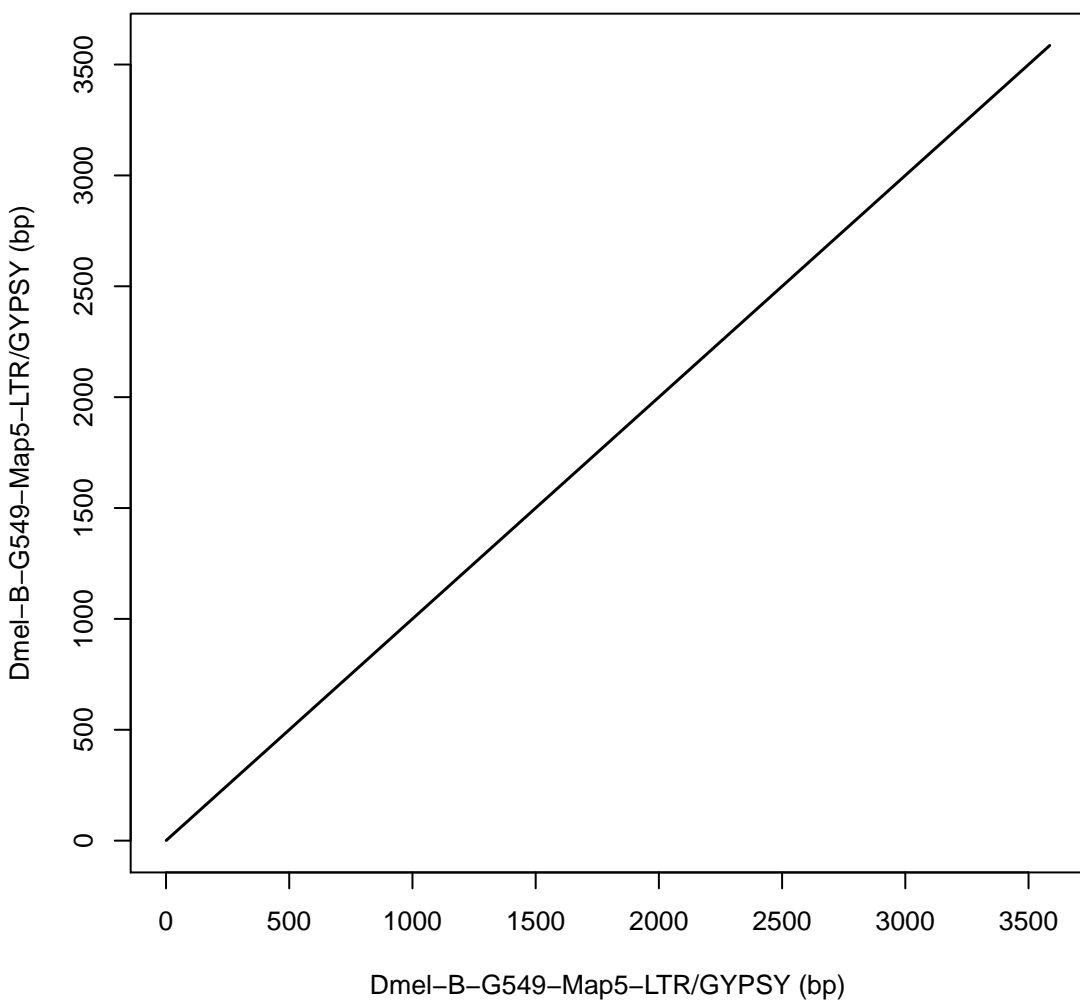
TE: Dmel-B-G549-Map5-LTR/GYPSY  
consensus size: 3586bp; fragments: 37; full length: 0 (>=3227.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

