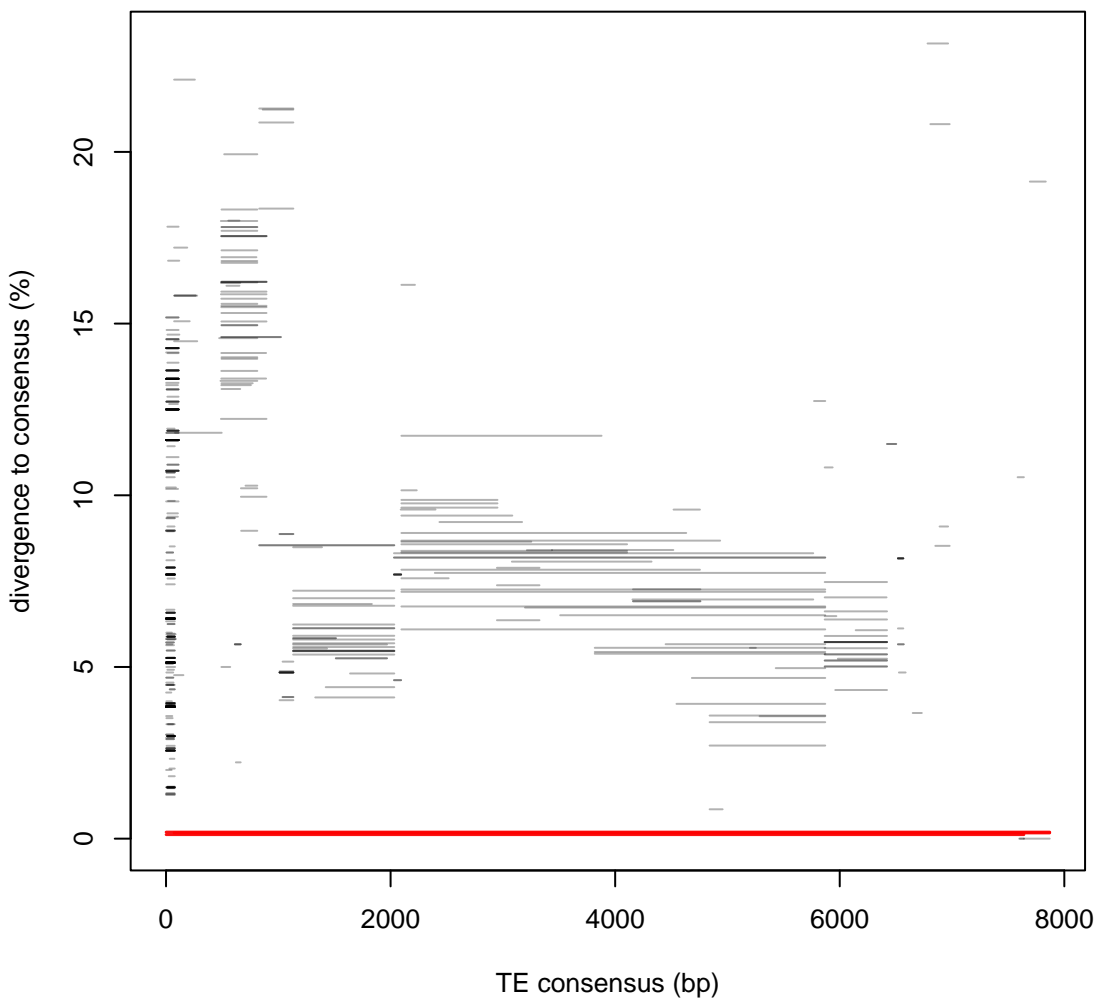
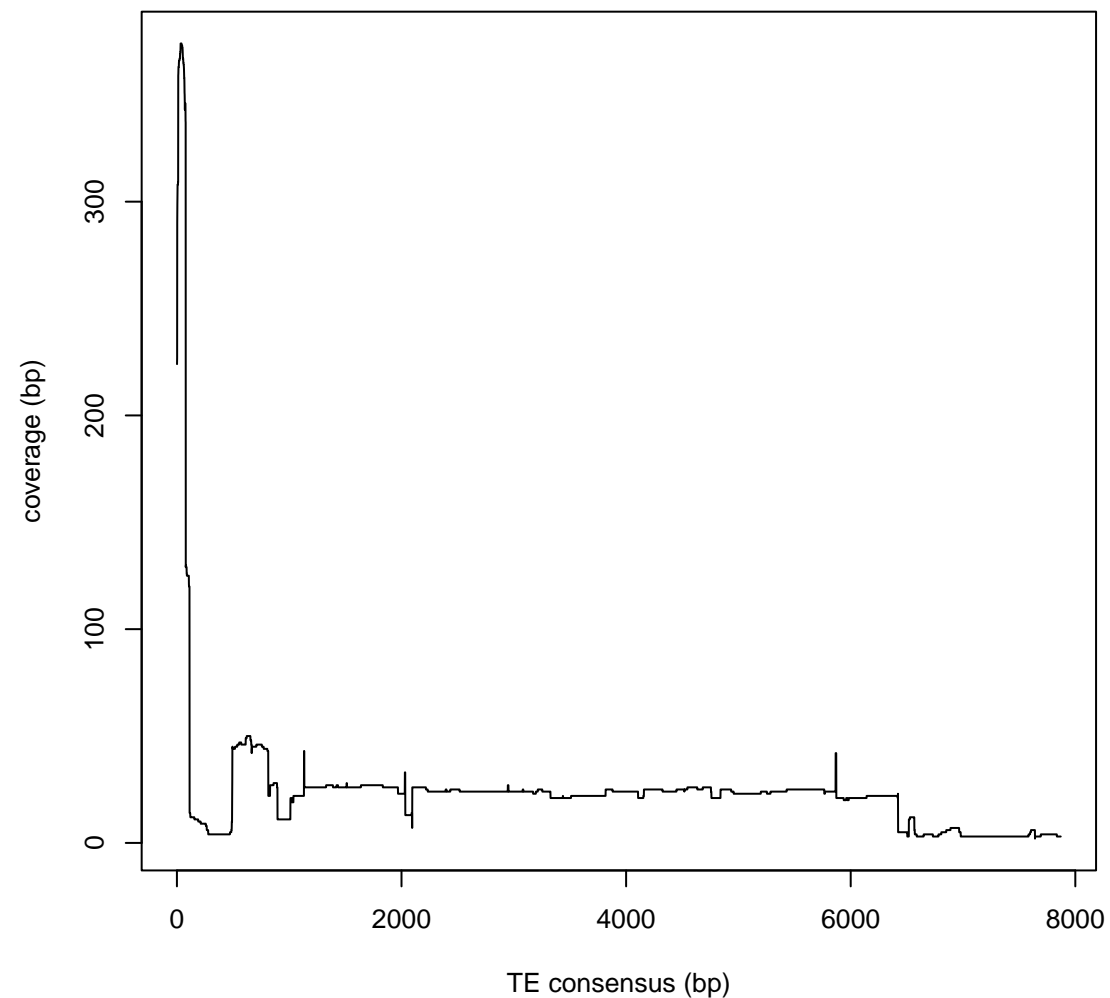


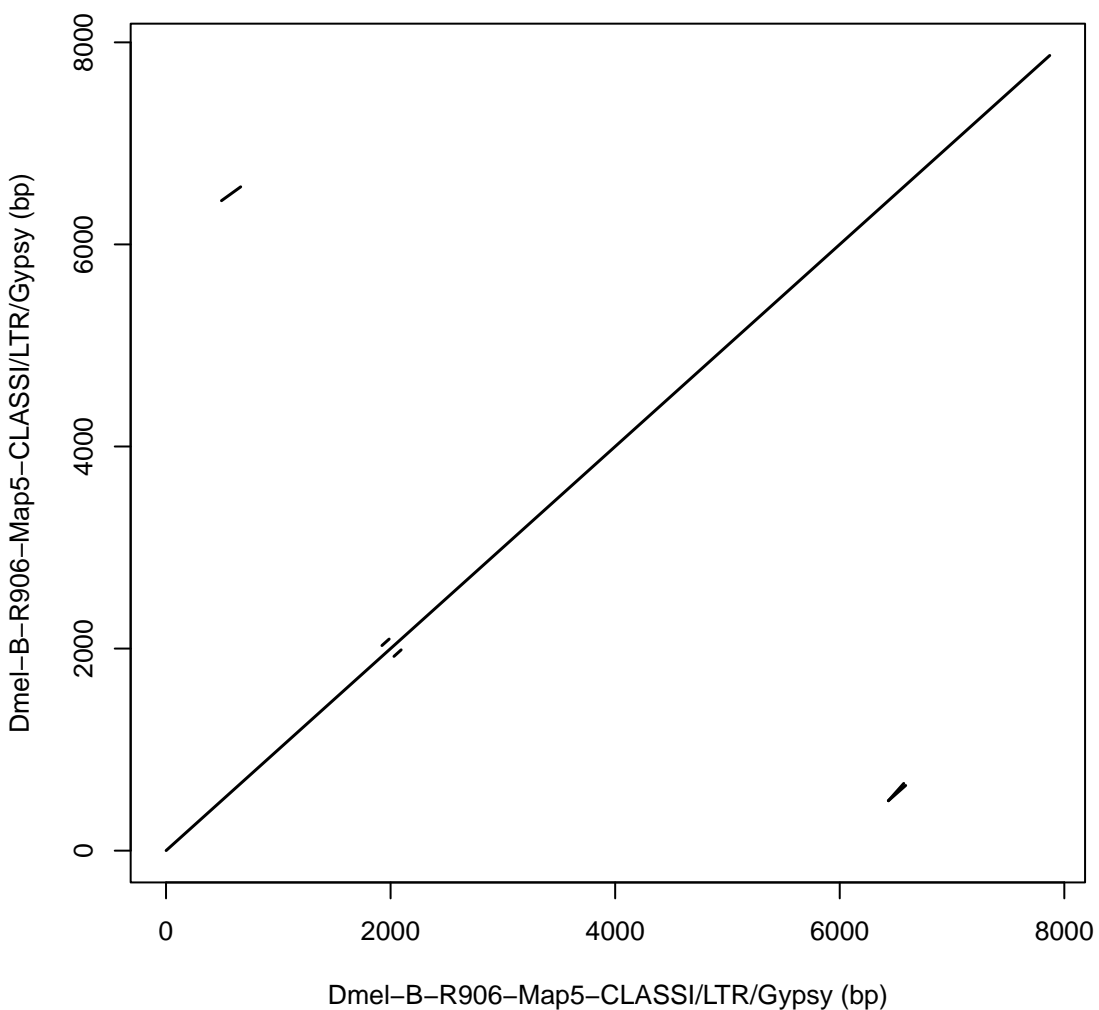
TE: Dmel-B-R906-Map5-CLASSI/LTR/Gypsy  
consensus size: 7870bp; fragments: 582; full length: 3 (>=7083bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

