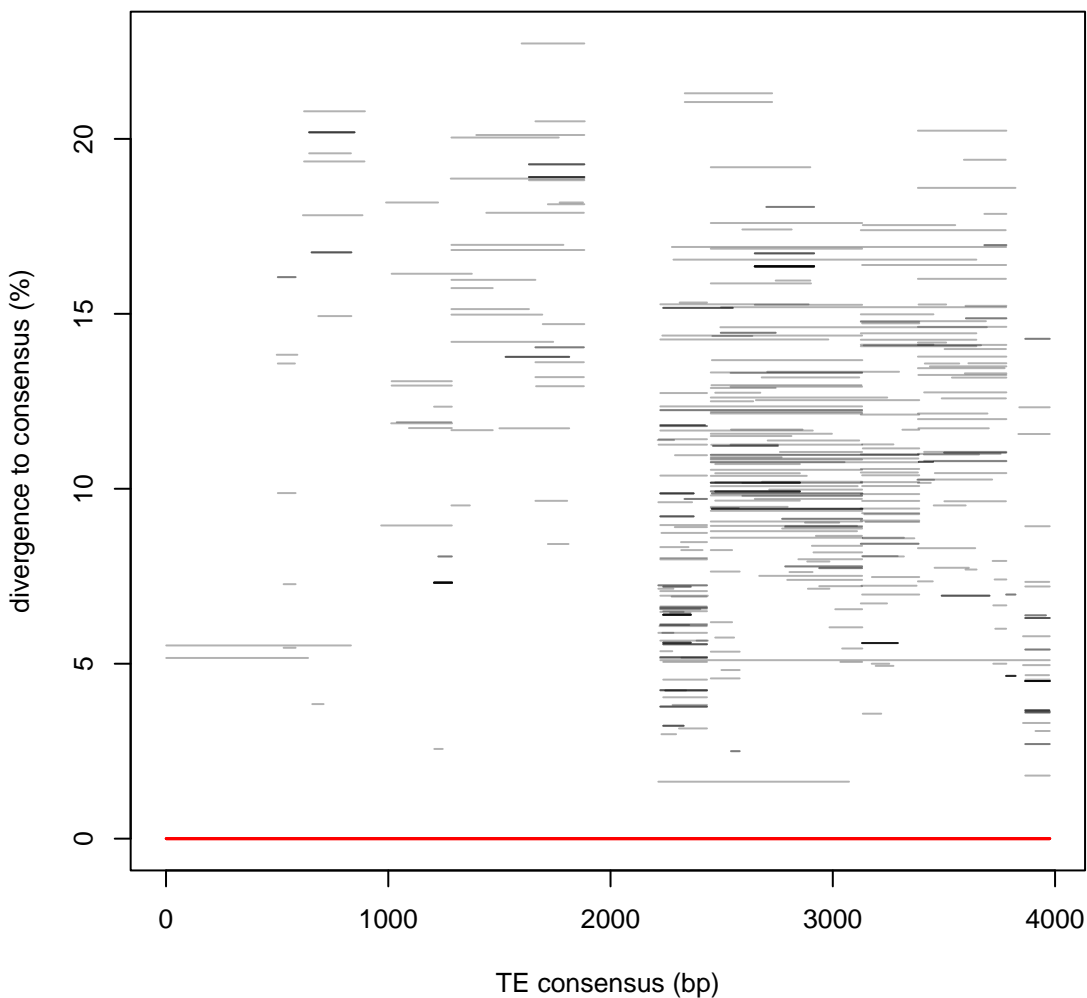
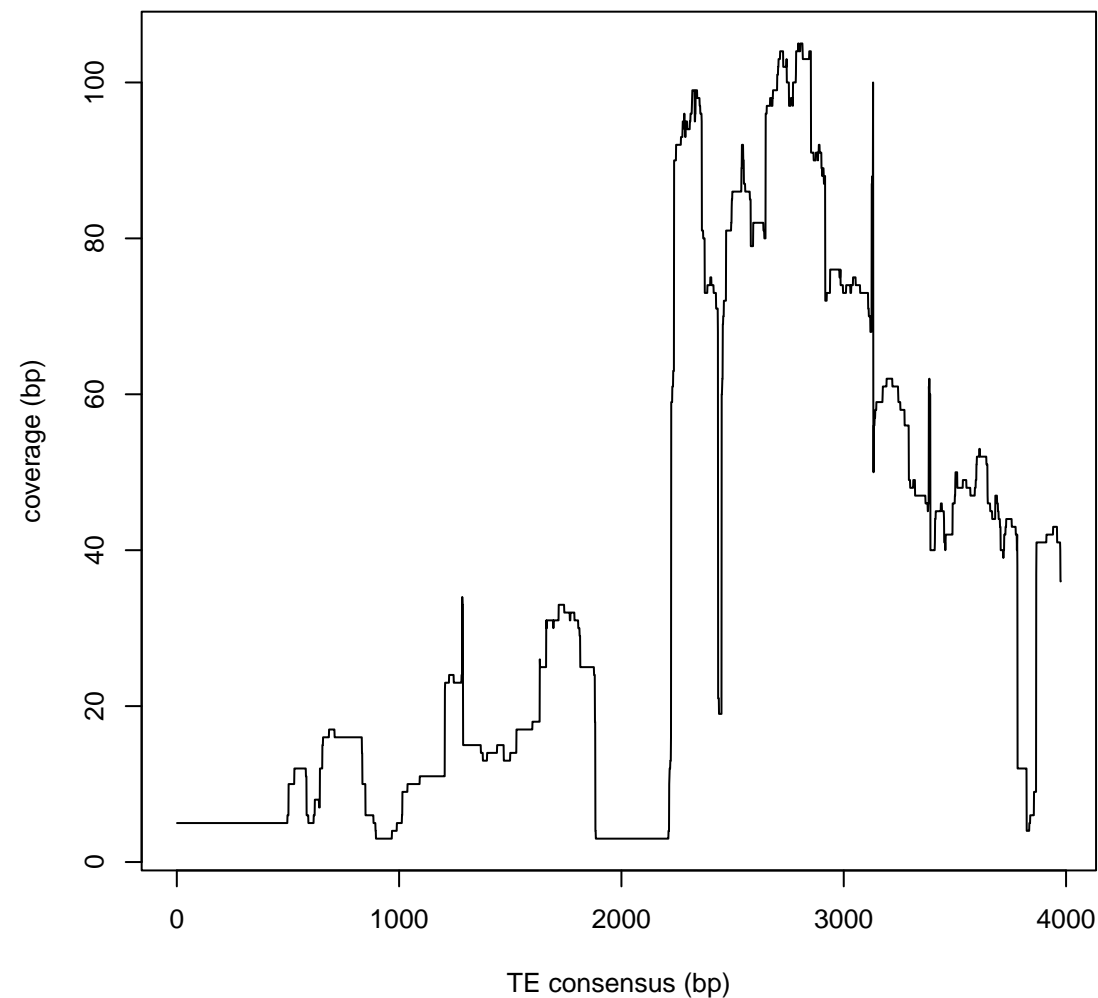


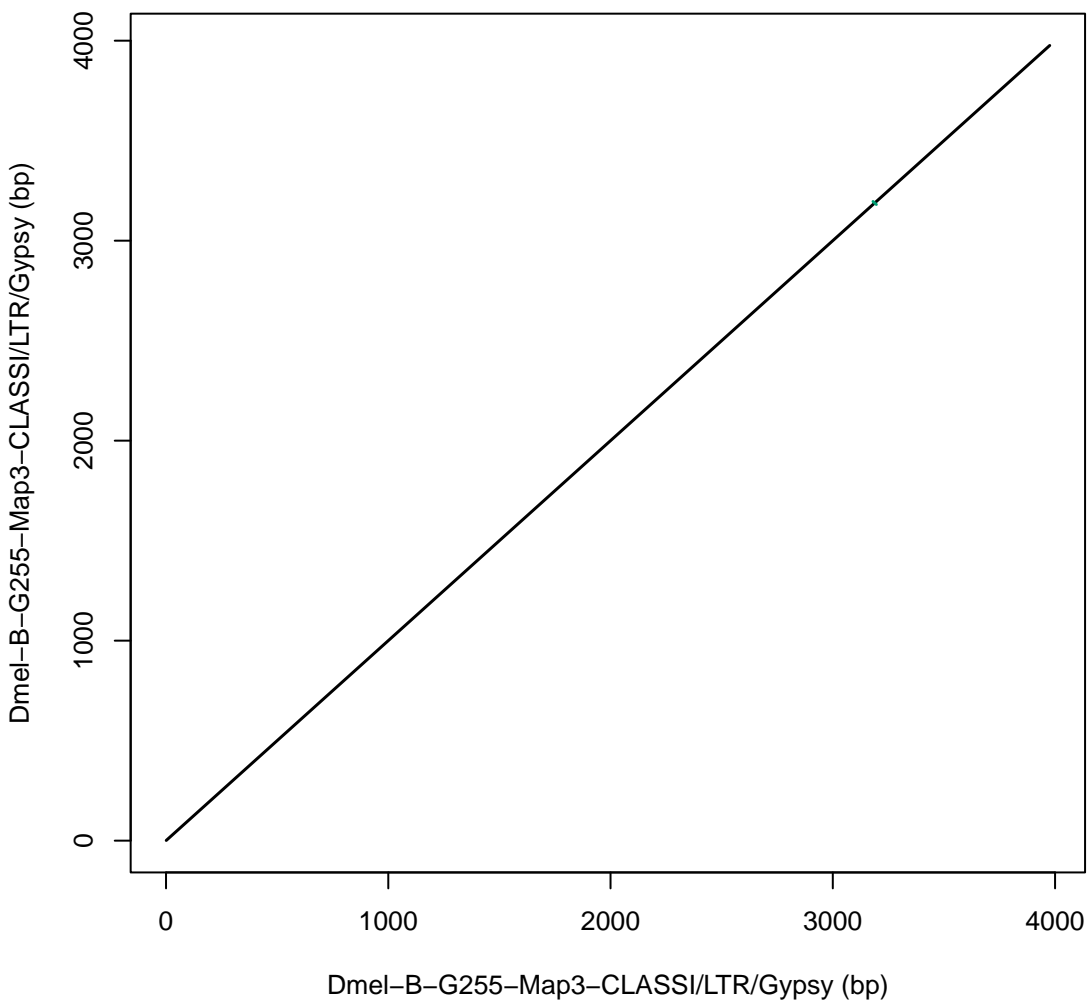
TE: Dmel-B-G255-Map3-CLASSI/LTR/Gypsy  
consensus size: 3976bp; fragments: 467; full length: 3 (>=3578.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

