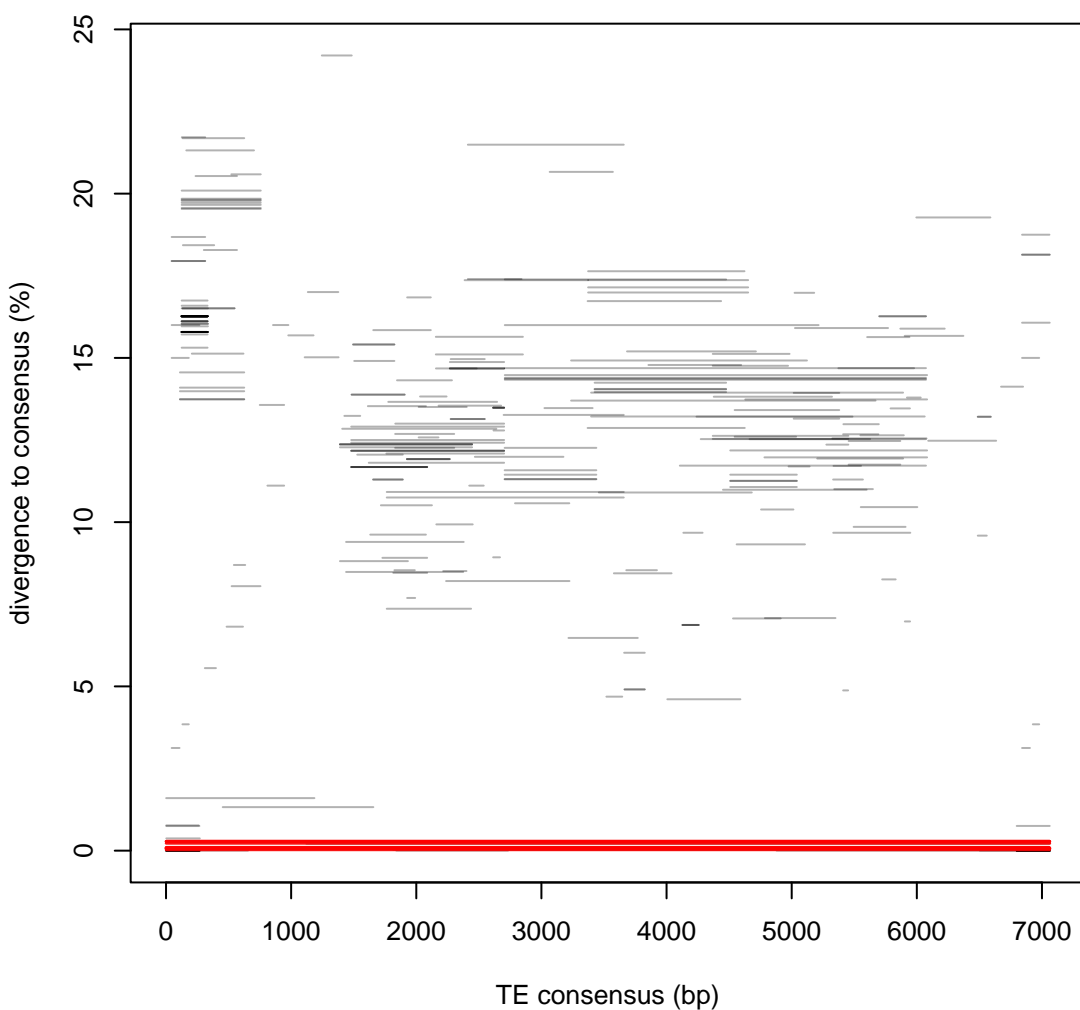
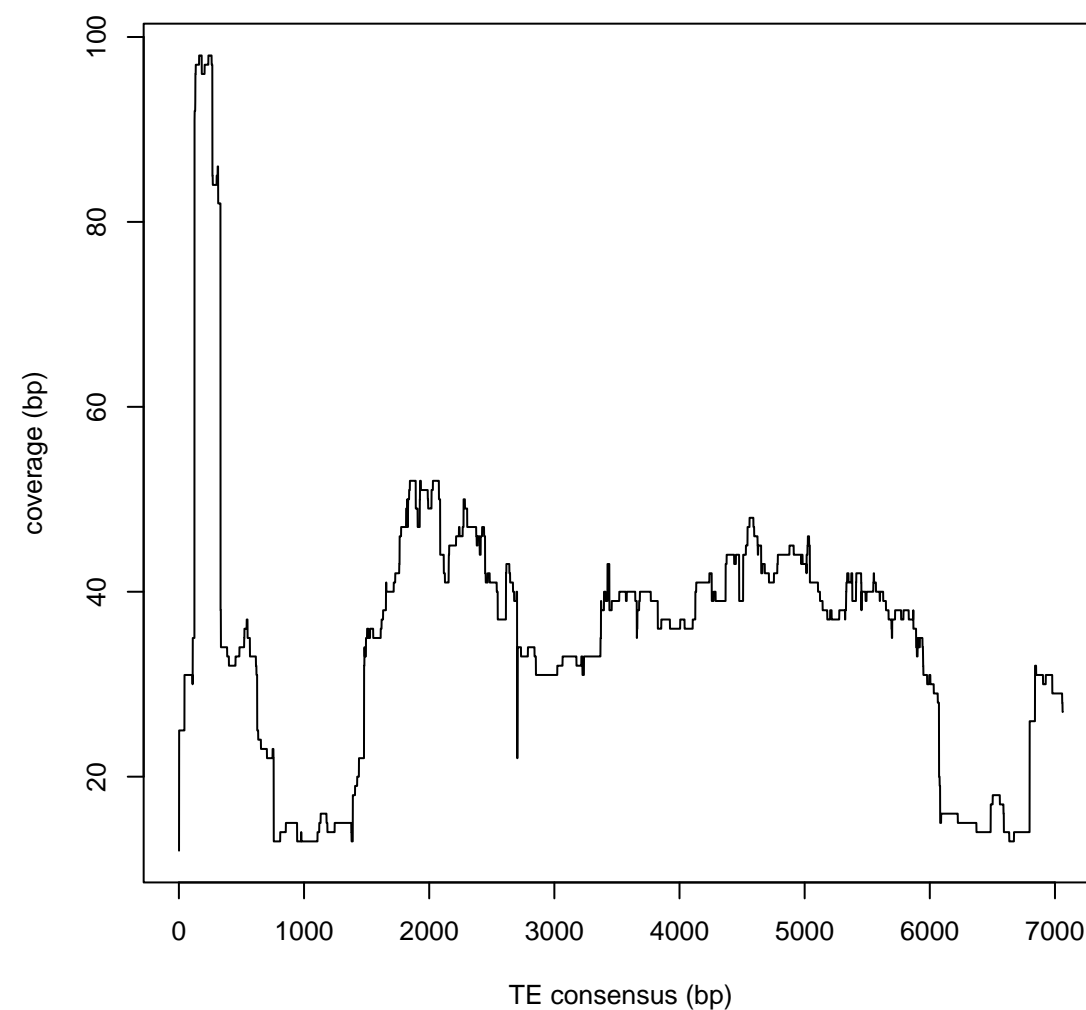


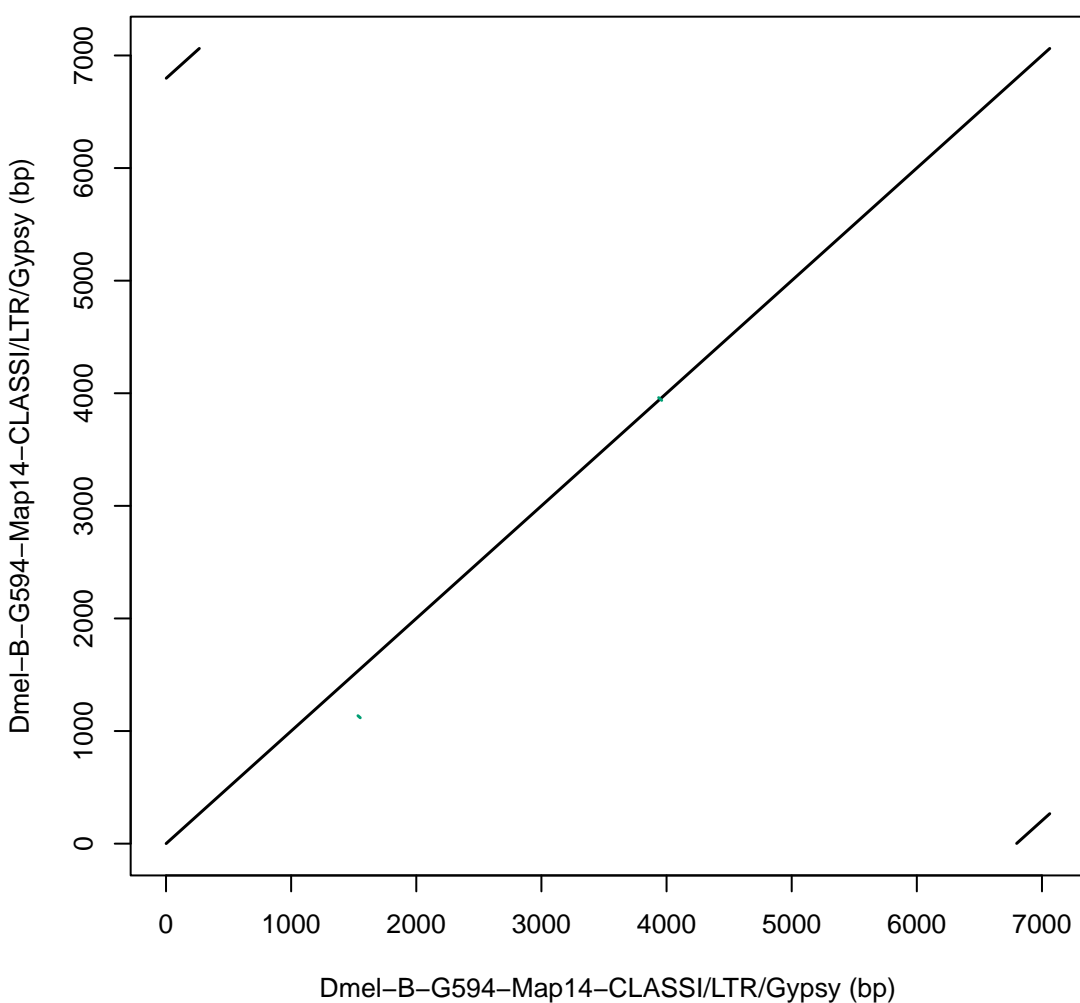
TE: Dmel-B-G594-Map14-CLASSI/LTR/Gypsy  
consensus size: 7062bp; fragments: 320; full length: 9 (>=6355.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

