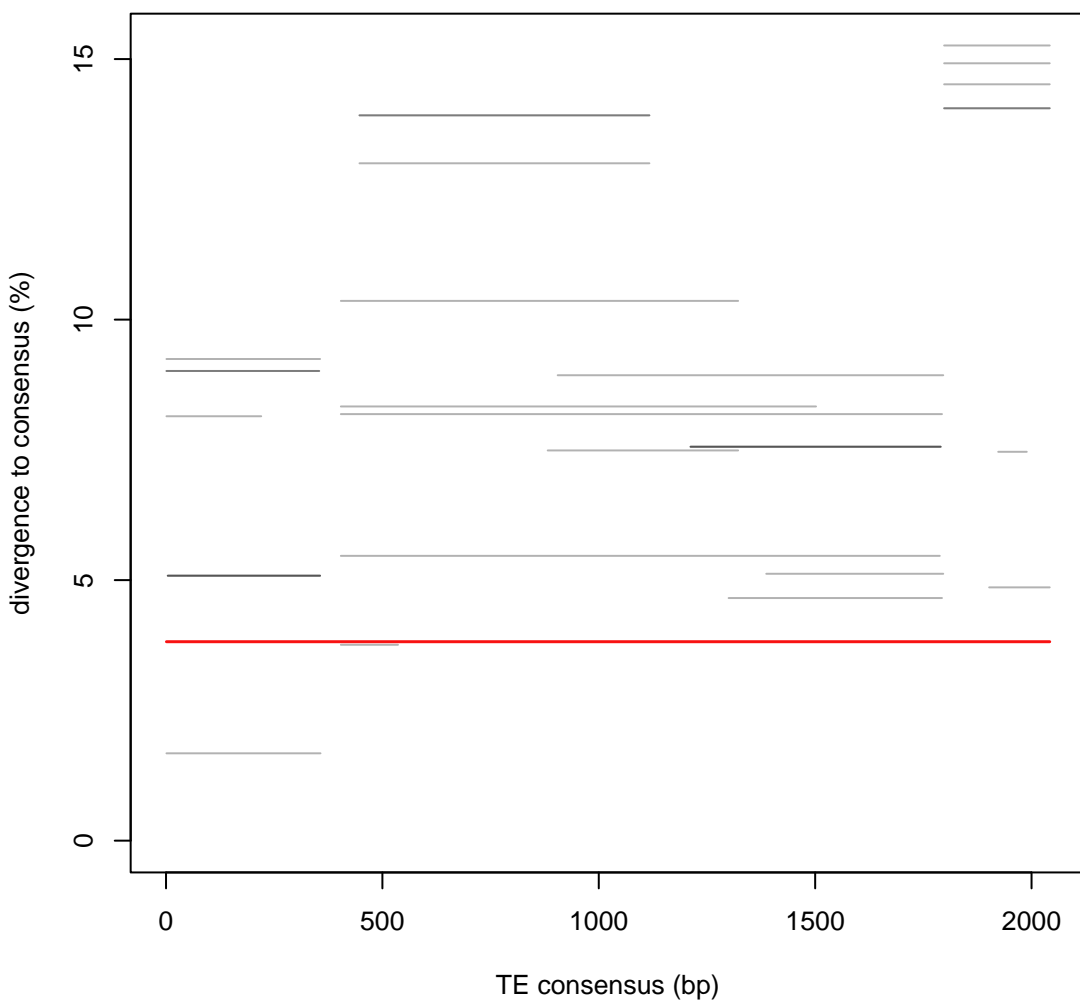
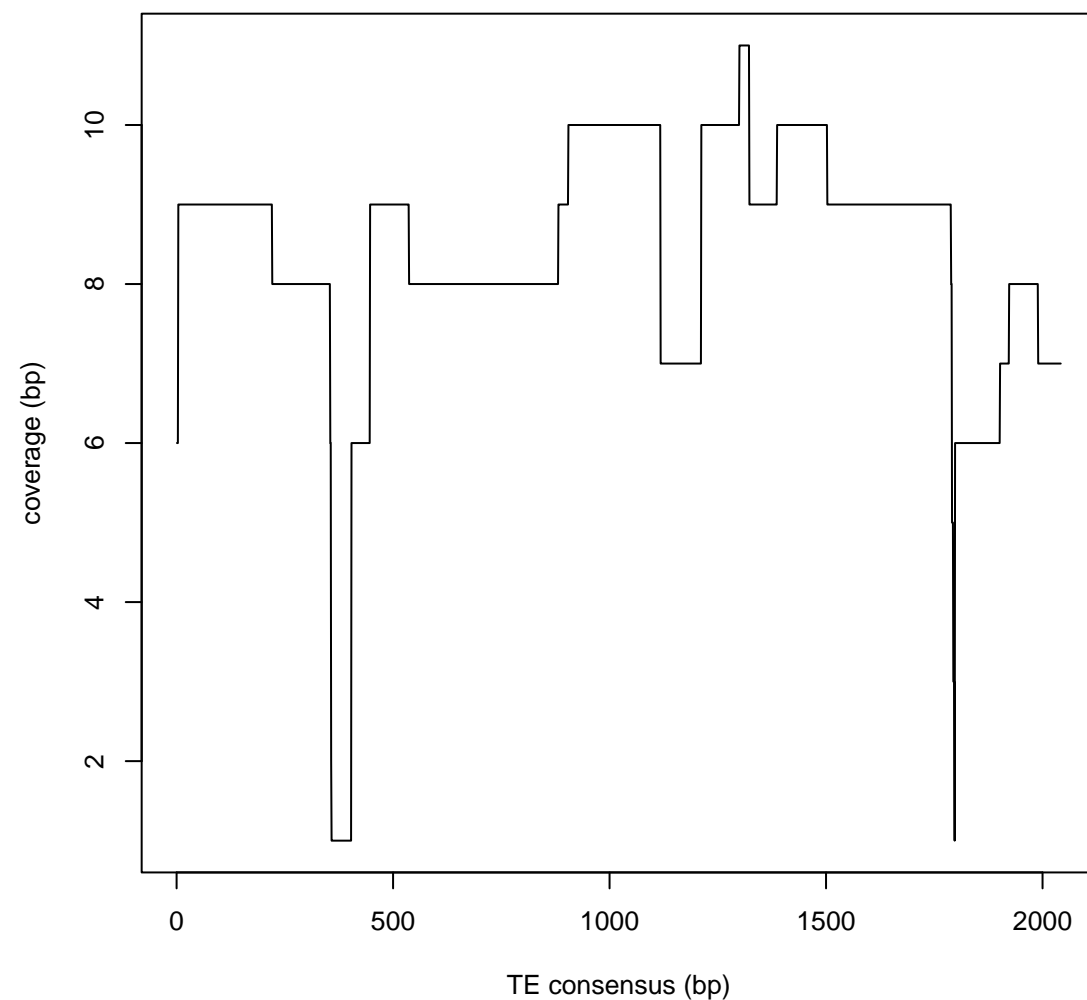


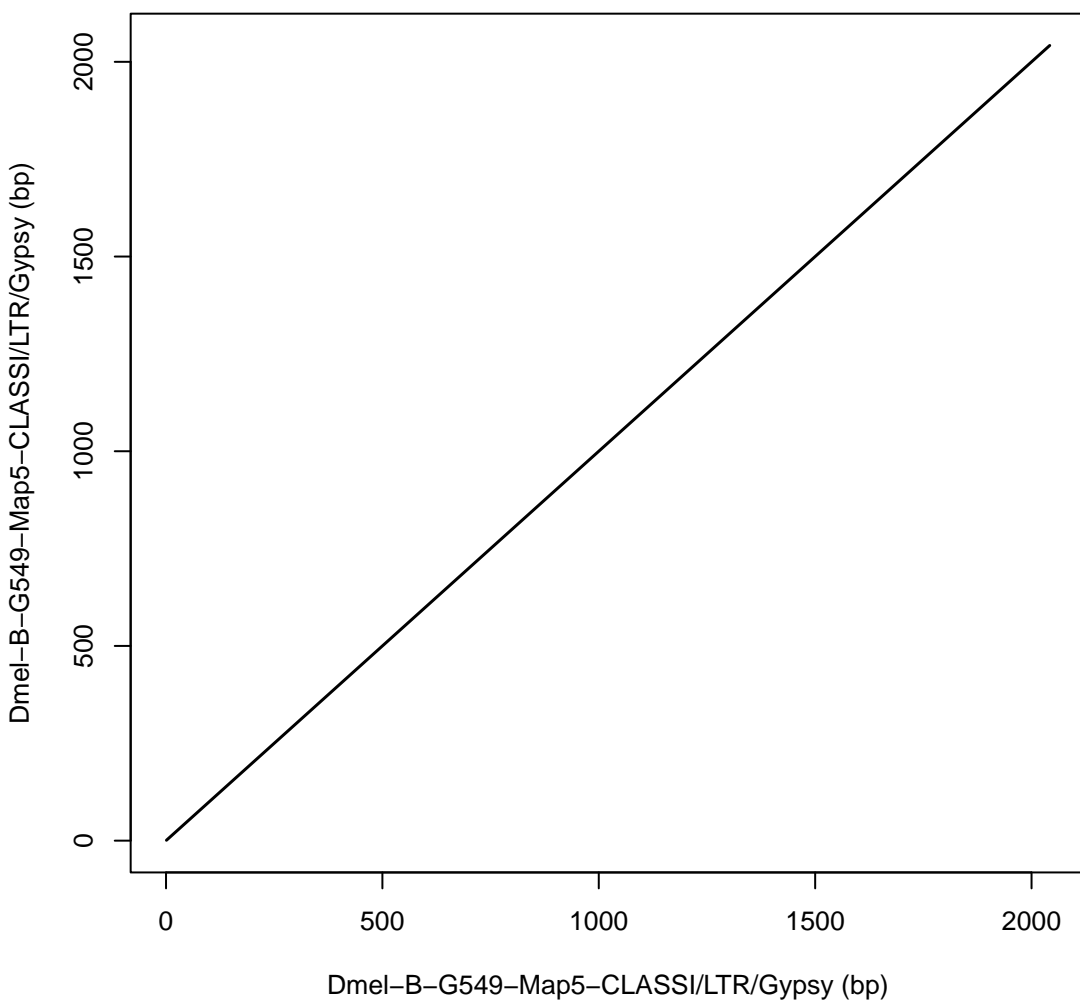
TE: Dmel-B-G549-Map5-CLASSI/LTR/Gypsy
consensus size: 2042bp; fragments: 31; full length: 1 (>=1837.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

