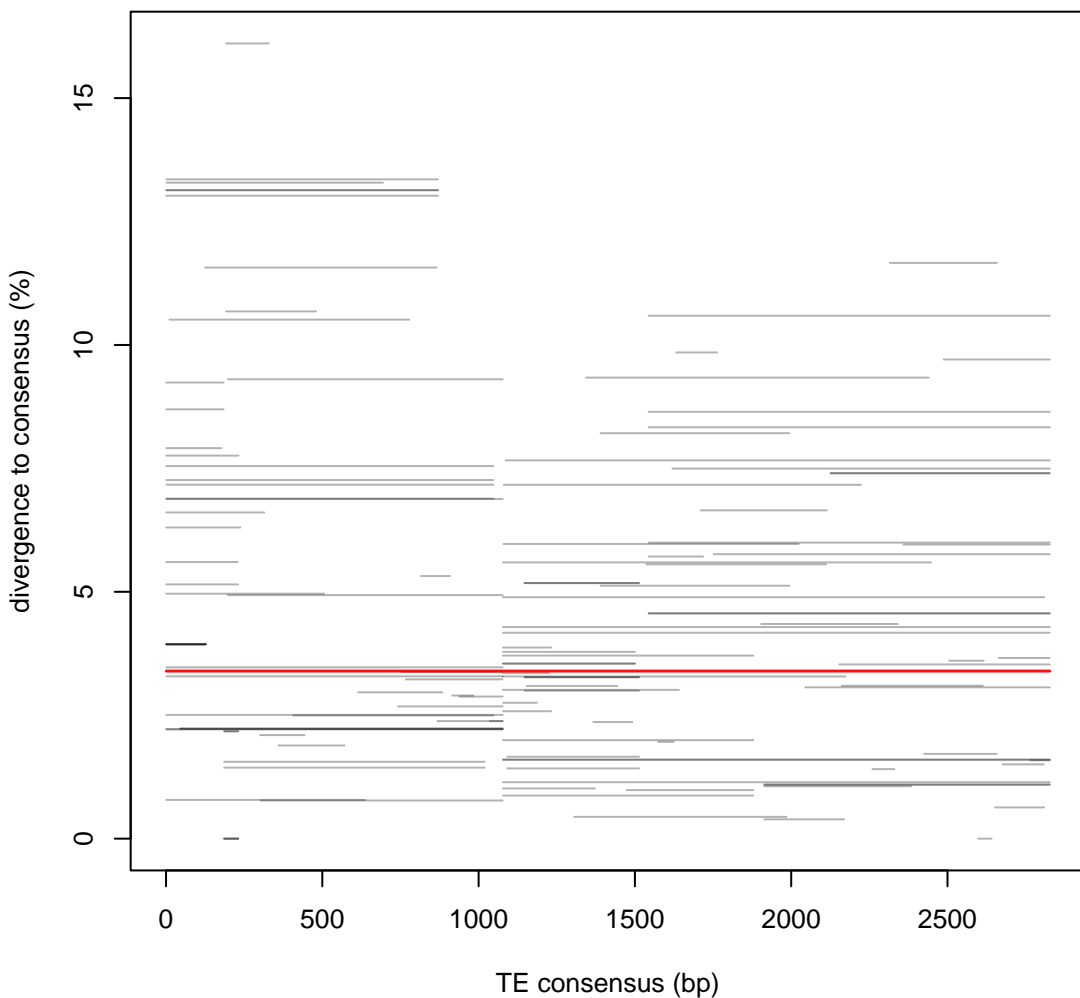
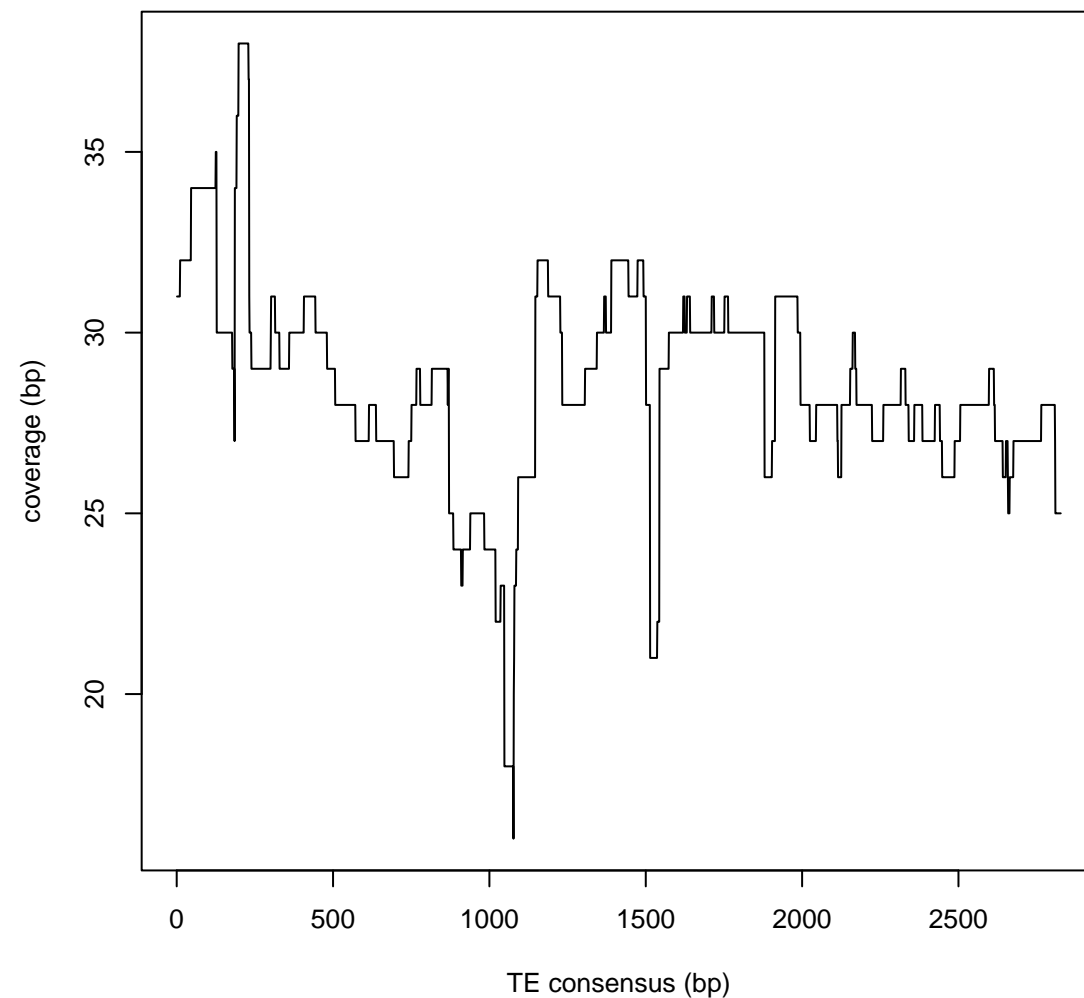


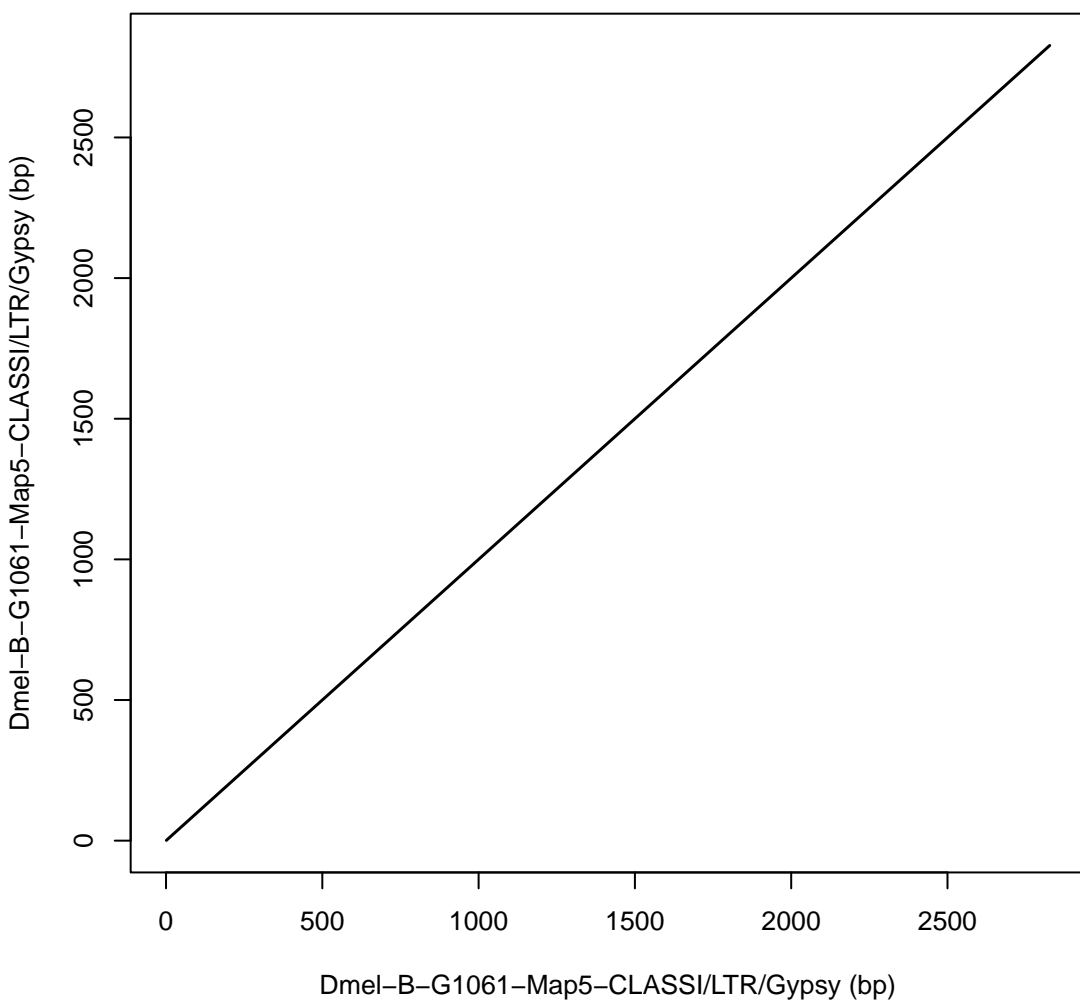
TE: Dmel-B-G1061-Map5-CLASSI/LTR/Gypsy  
consensus size: 2827bp; fragments: 130; full length: 1 (>=2544.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

