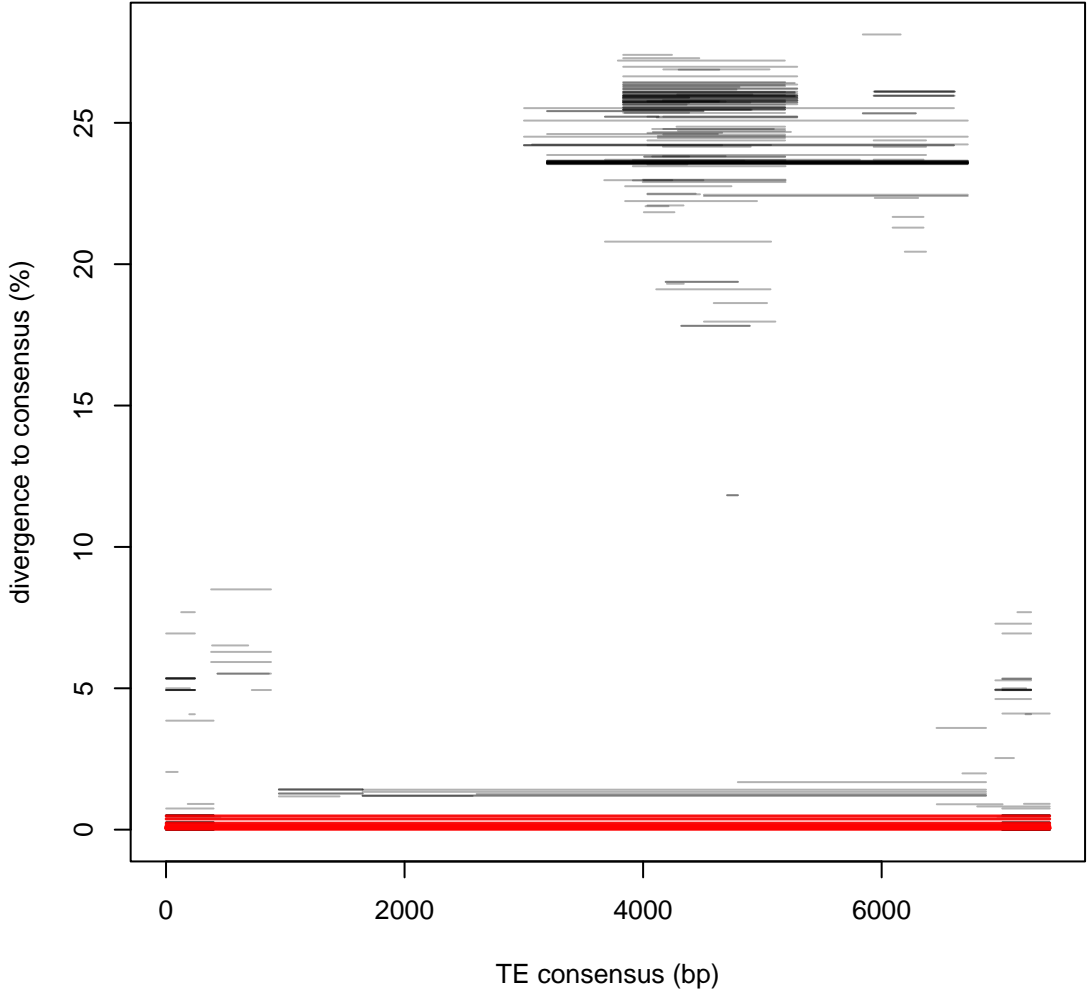
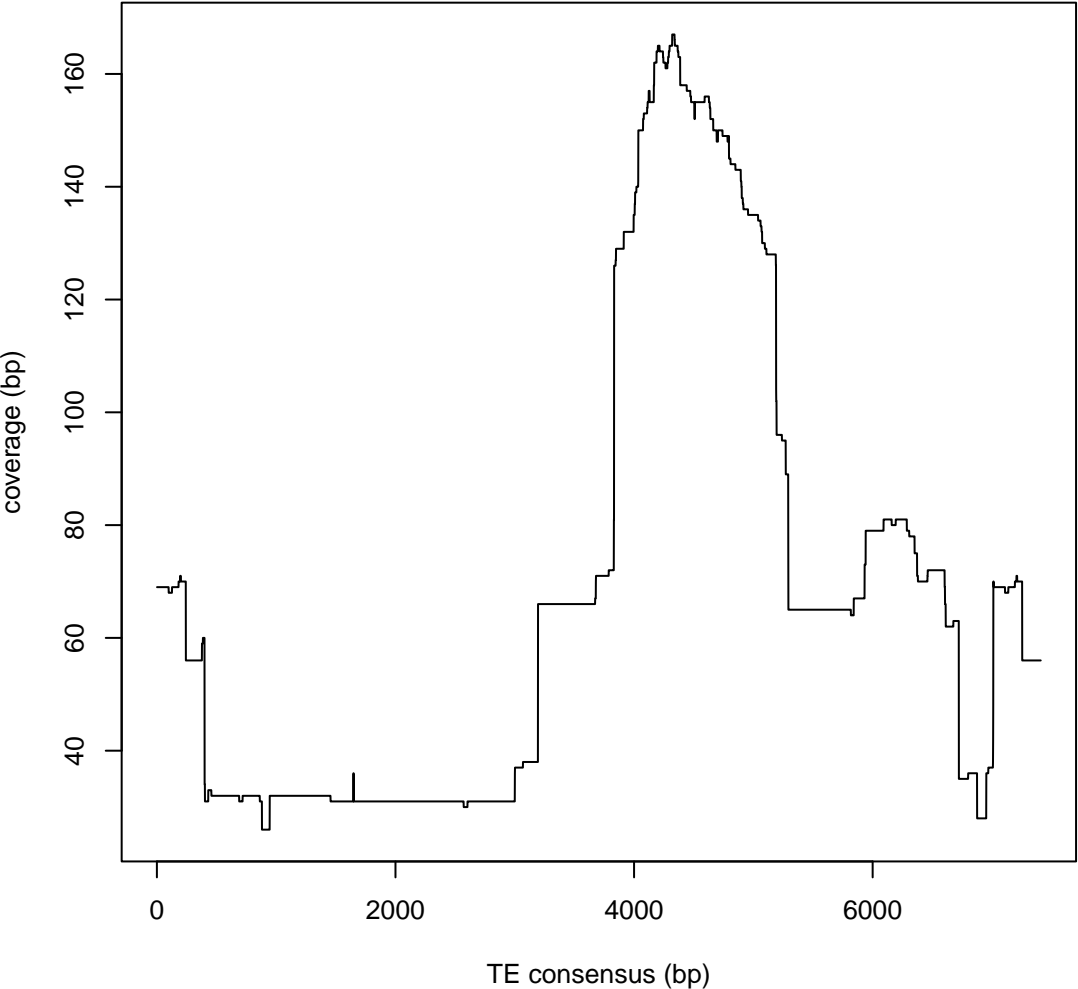


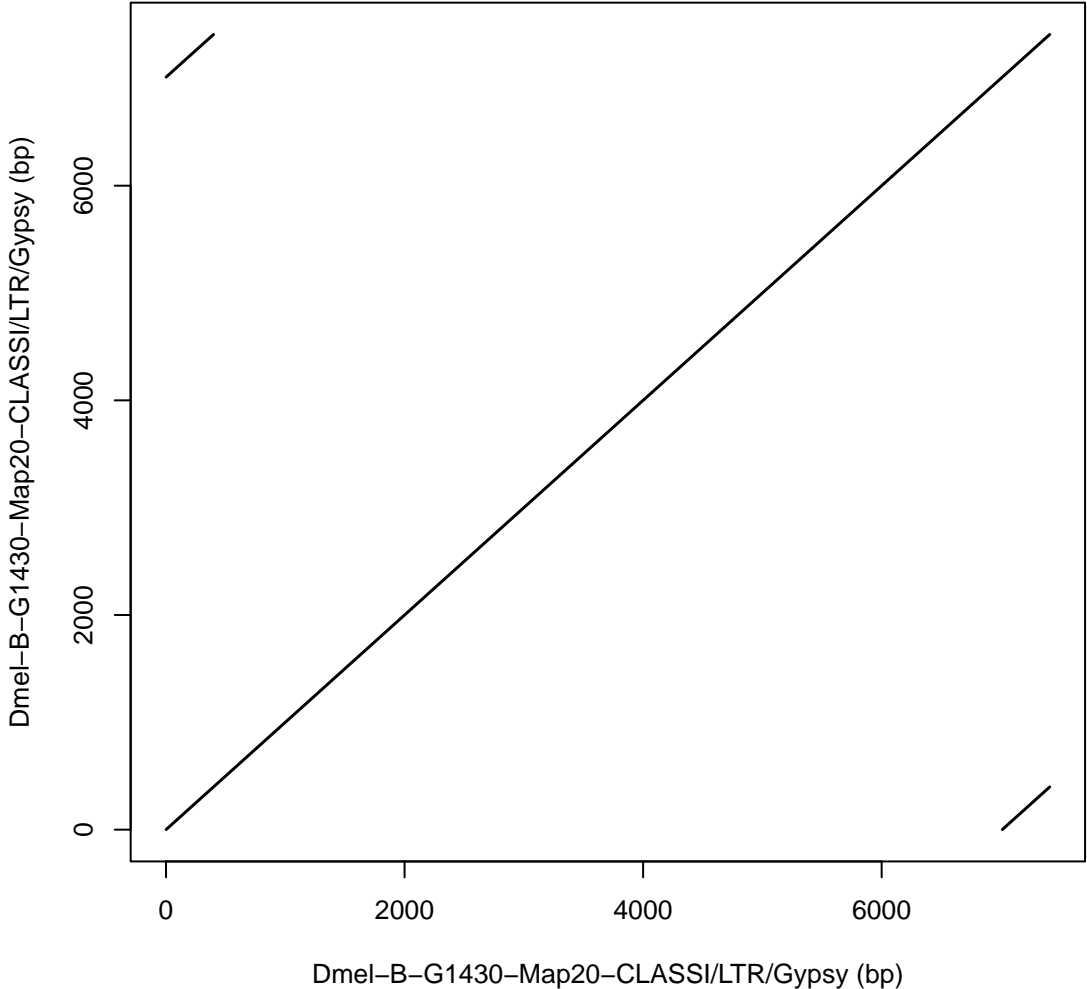
TE: Dmel-B-G1430-Map20-CLASSI/LTR/Gypsy  
consensus size: 7409bp; fragments: 307; full length: 26 (>=6668.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

