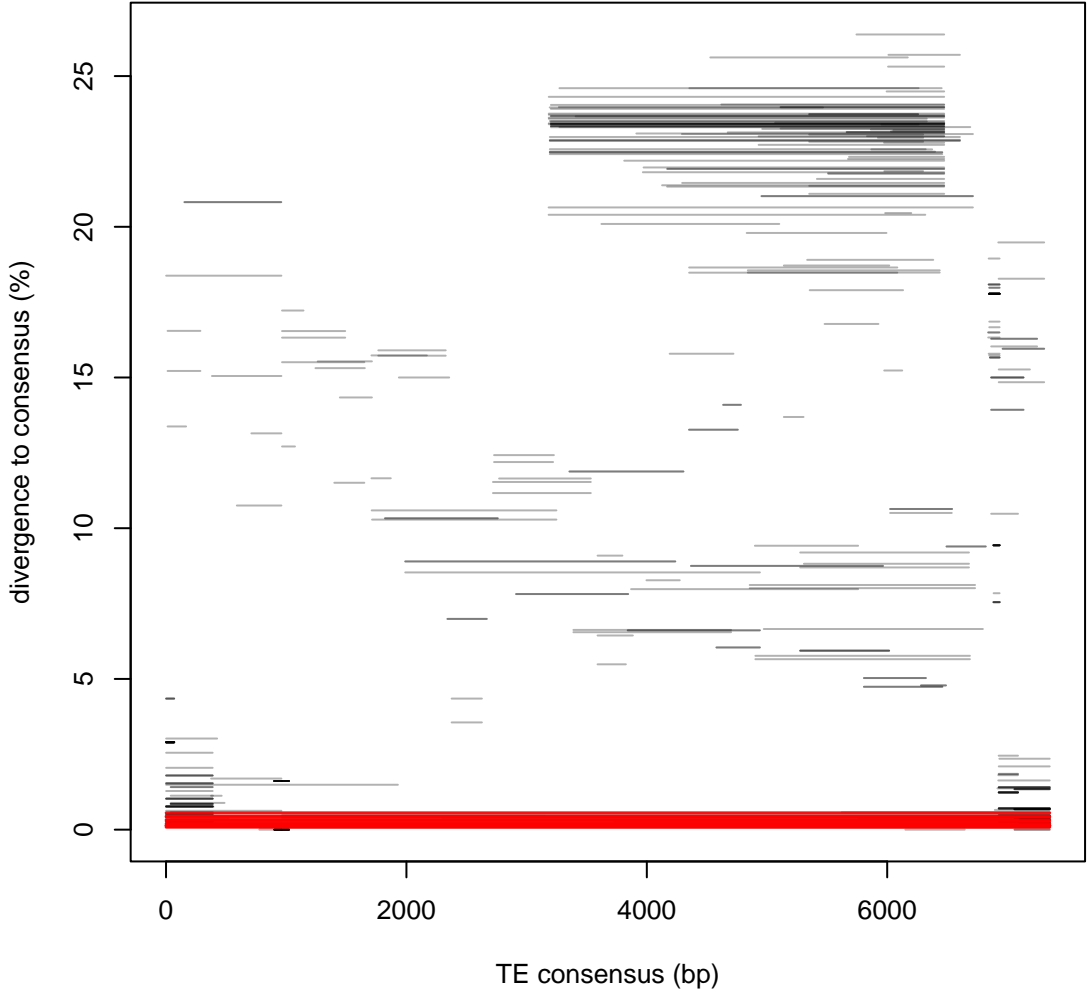
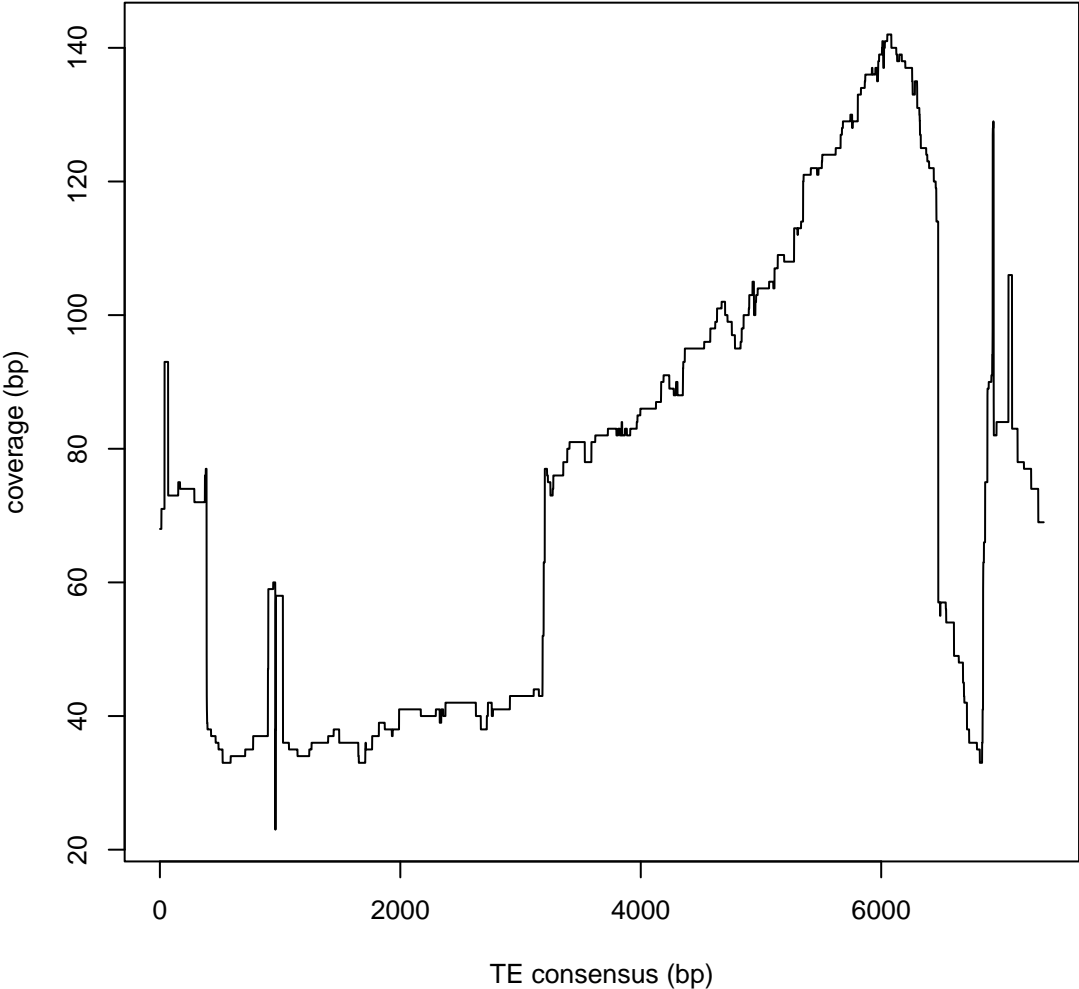


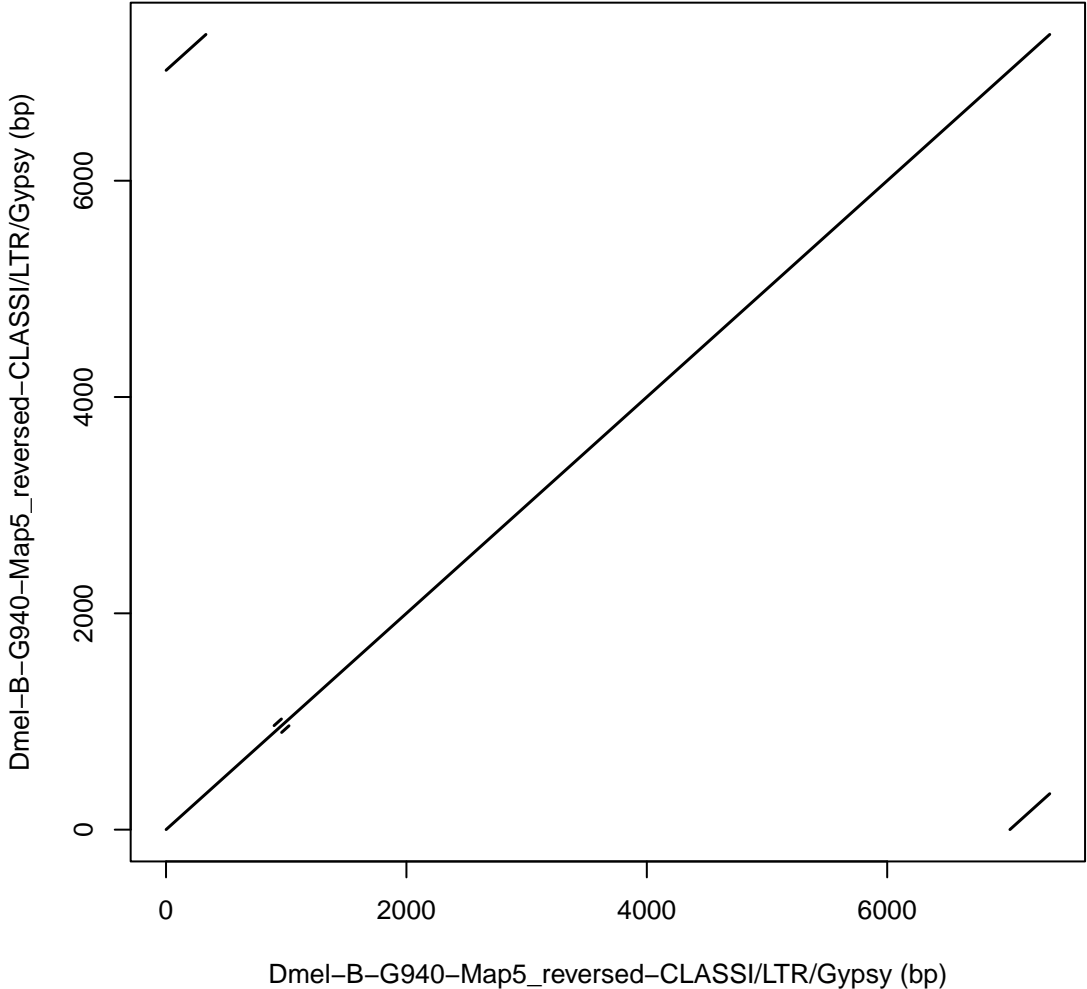
TE: Dmel-B-G940-Map5_reversed-CLASSI/LTR/Gypsy
consensus size: 7352bp; fragments: 458; full length: 20 (>=6616.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

