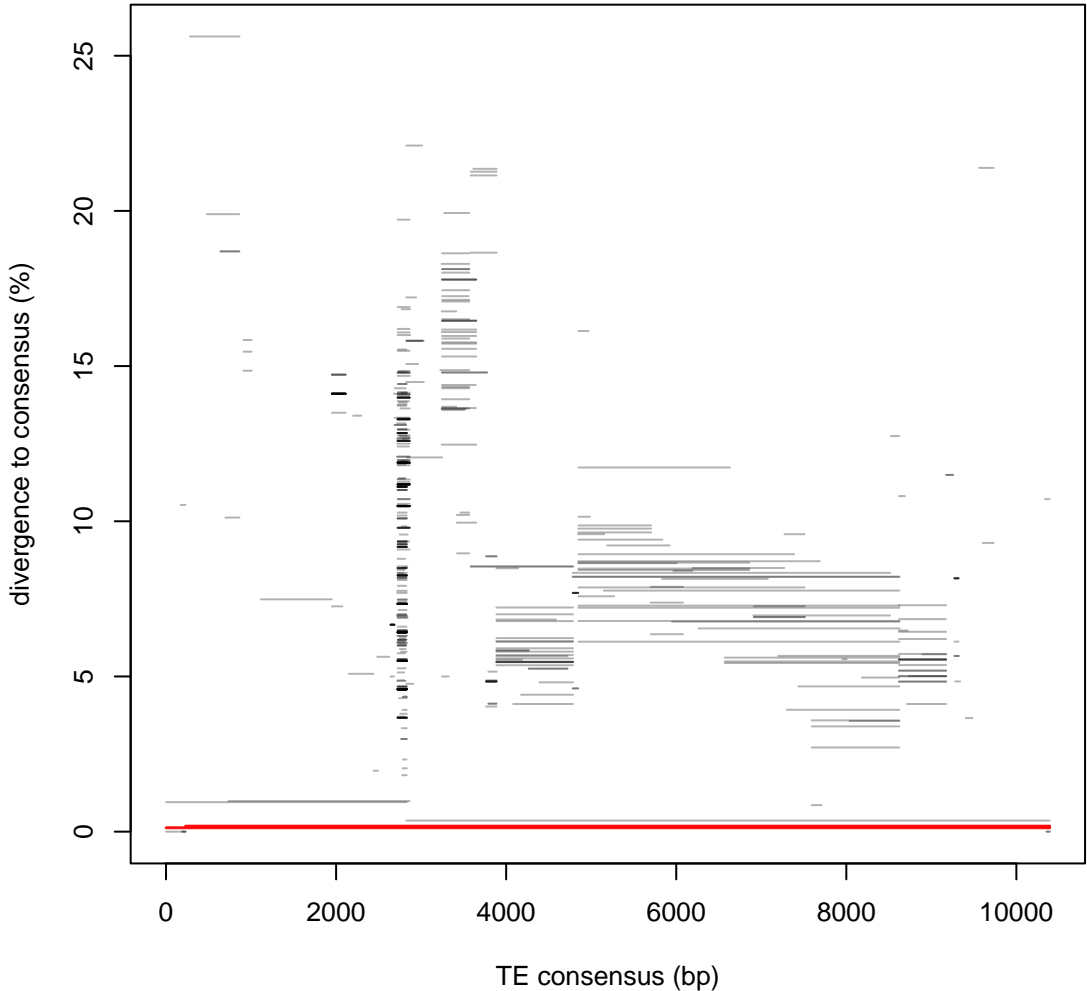
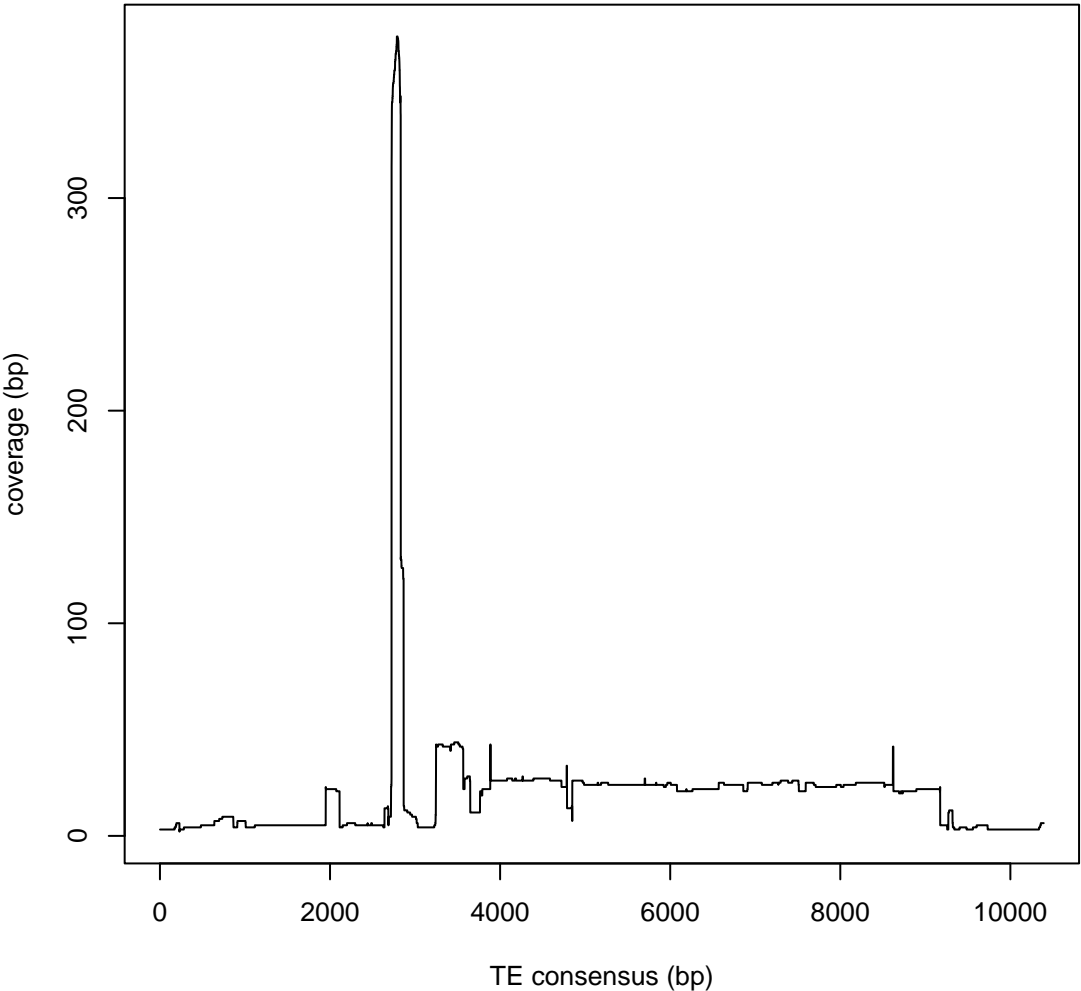


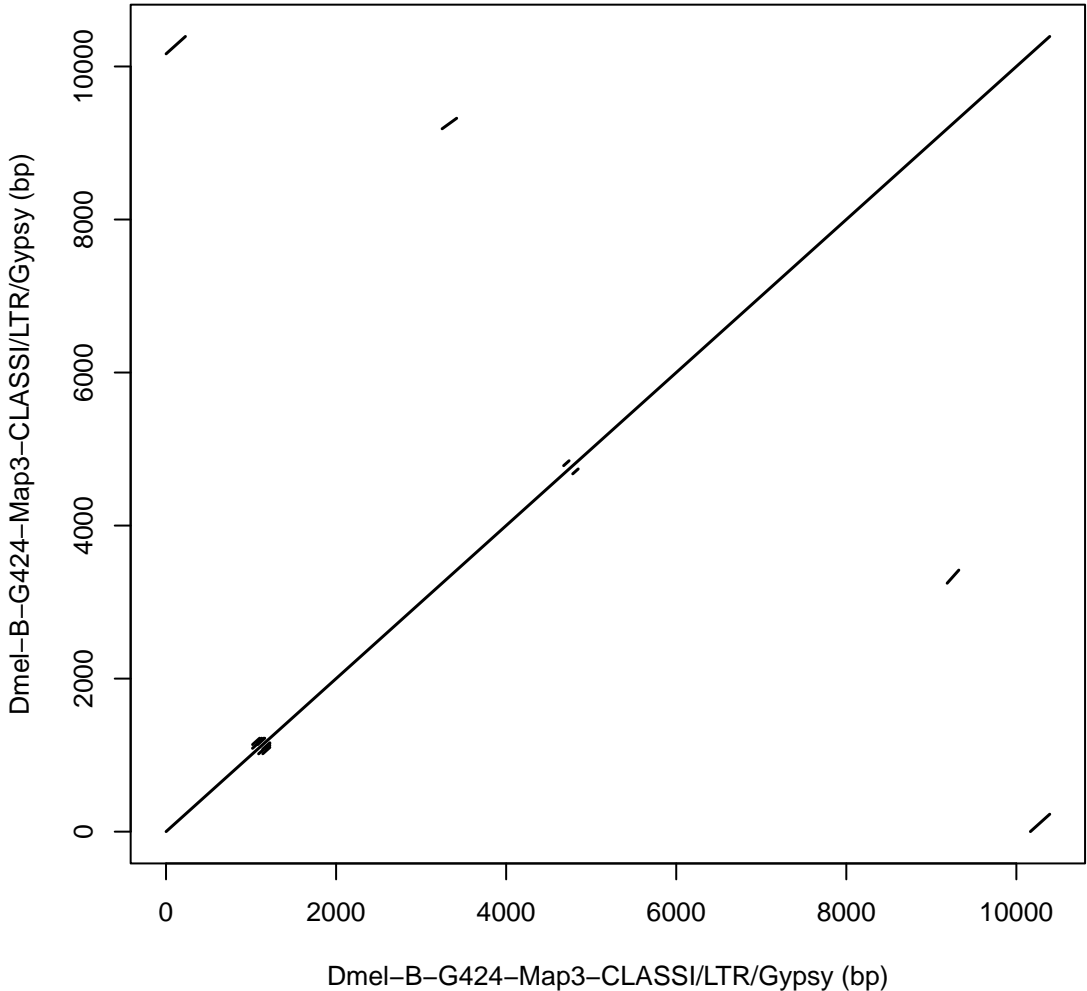
TE: Dmel-B-G424-Map3-CLASSI/LTR/Gypsy
consensus size: 10392bp; fragments: 616; full length: 2 (>=9352.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

