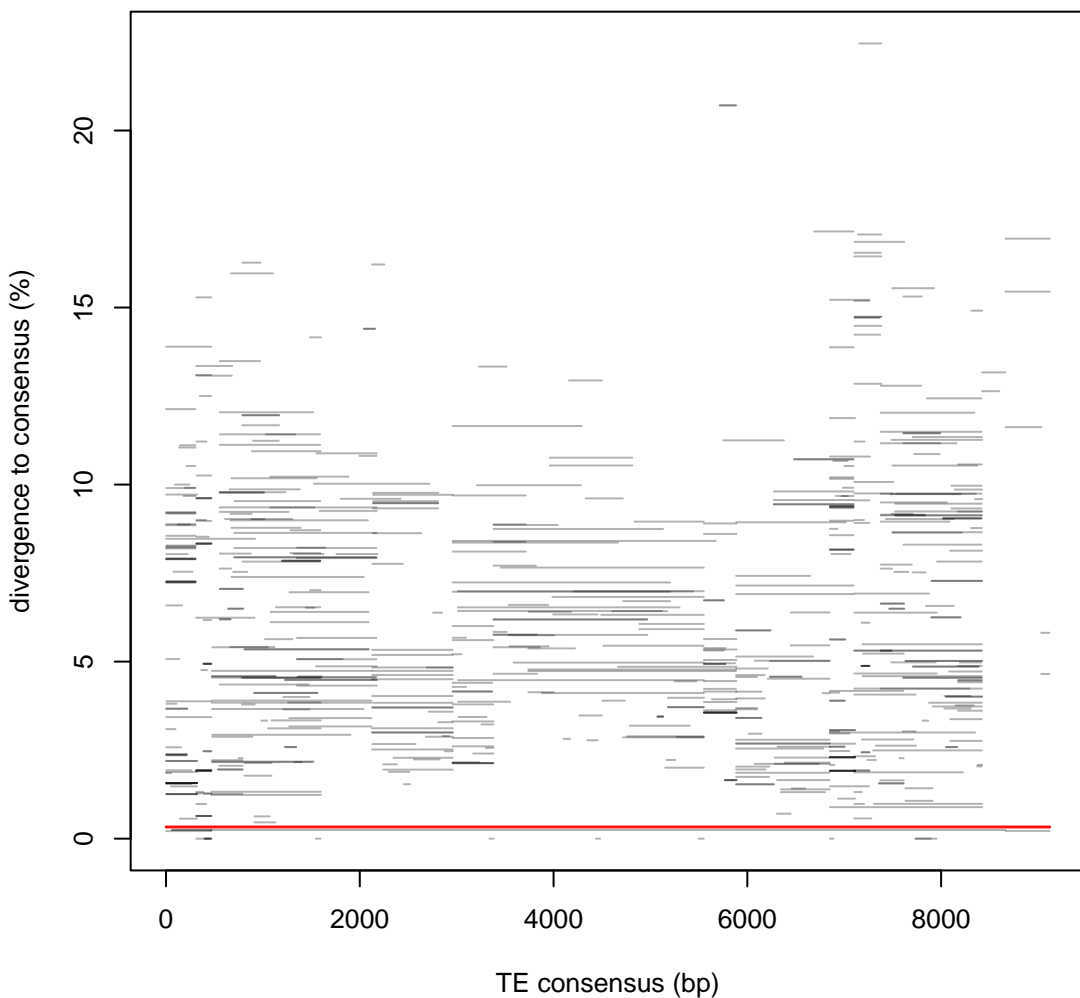
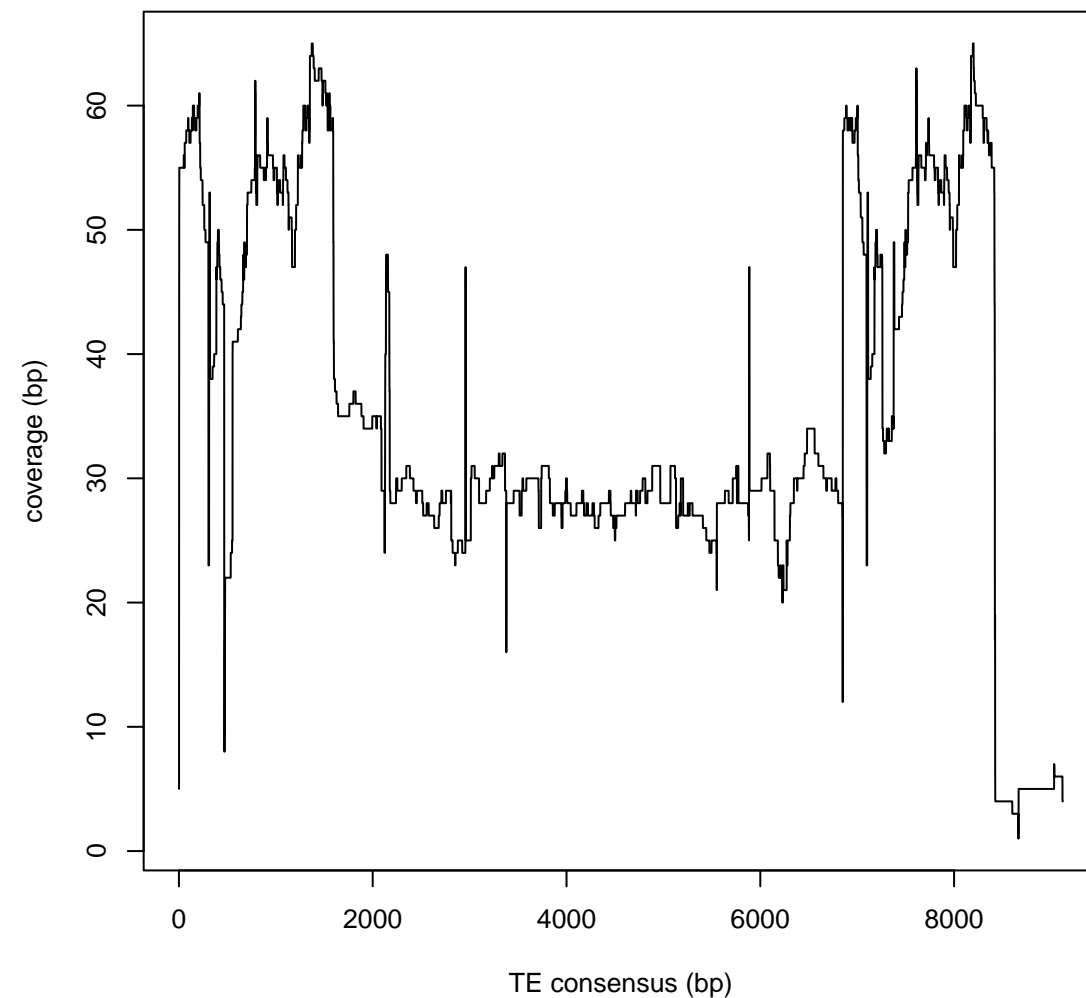


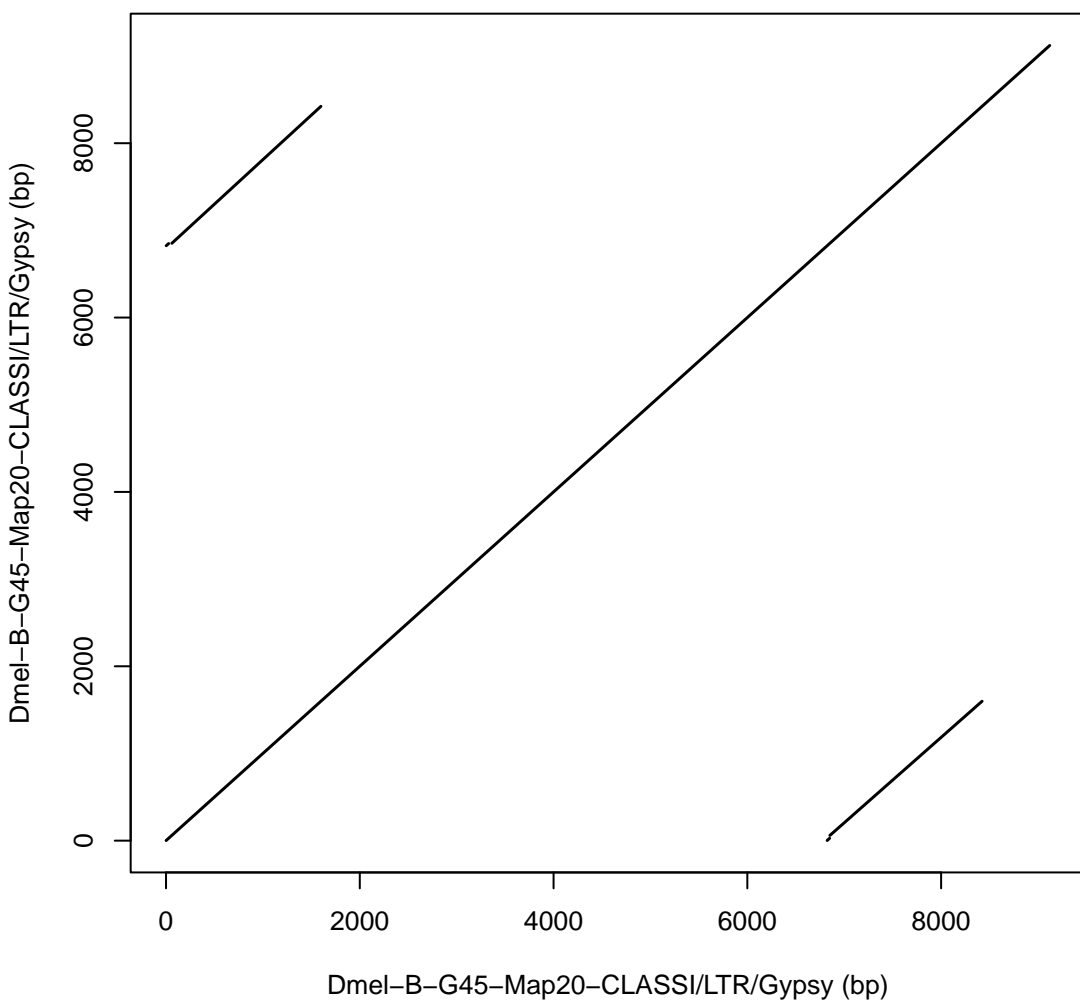
TE: Dmel-B-G45-Map20-CLASSI/LTR/Gypsy
consensus size: 9121bp; fragments: 645; full length: 1 (>=8208.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

