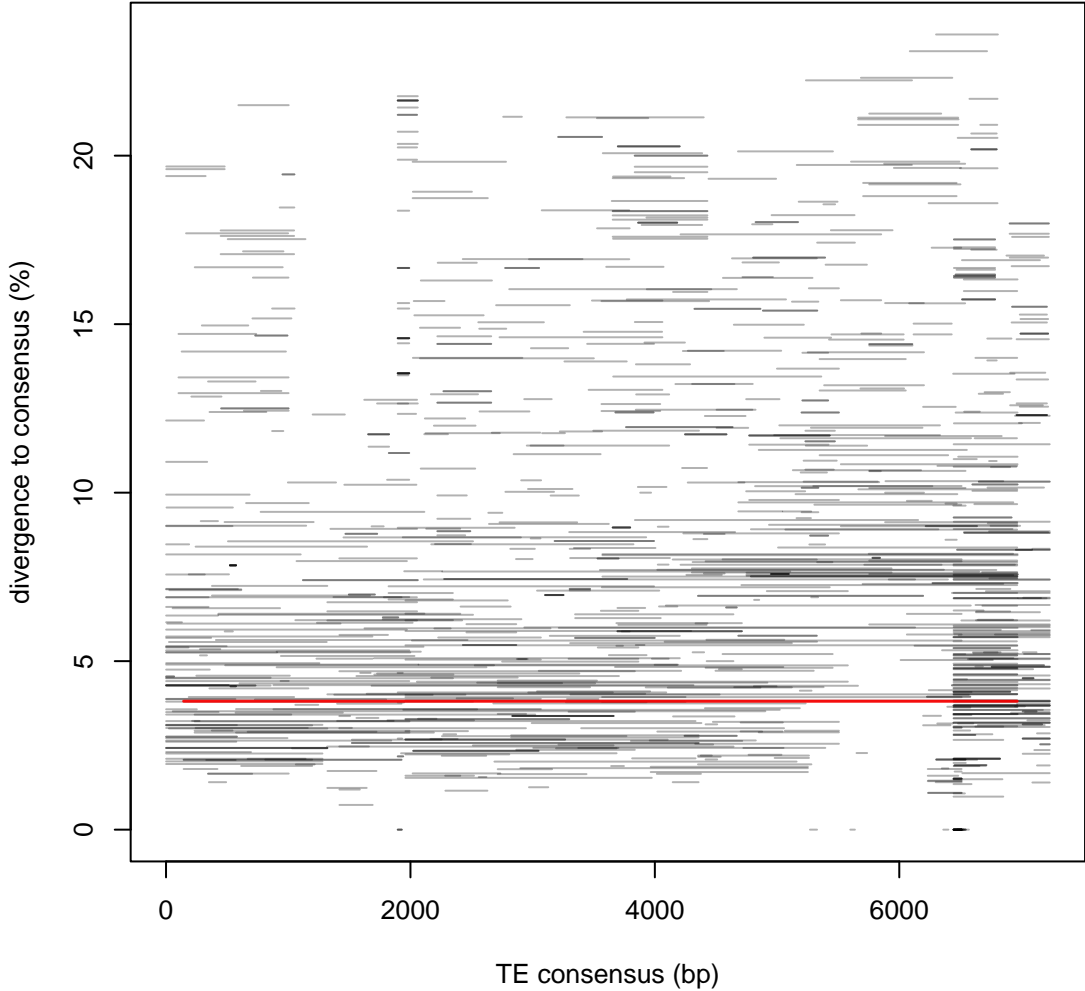
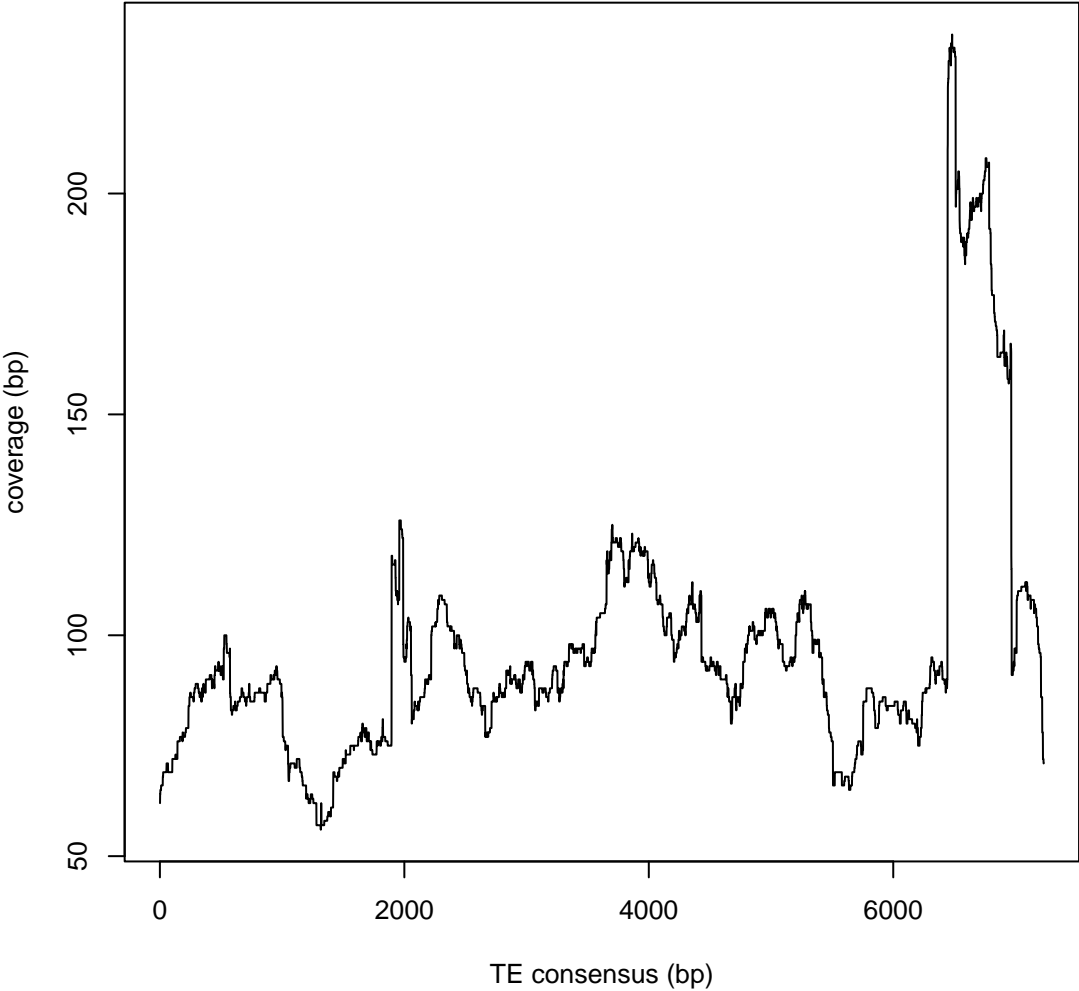


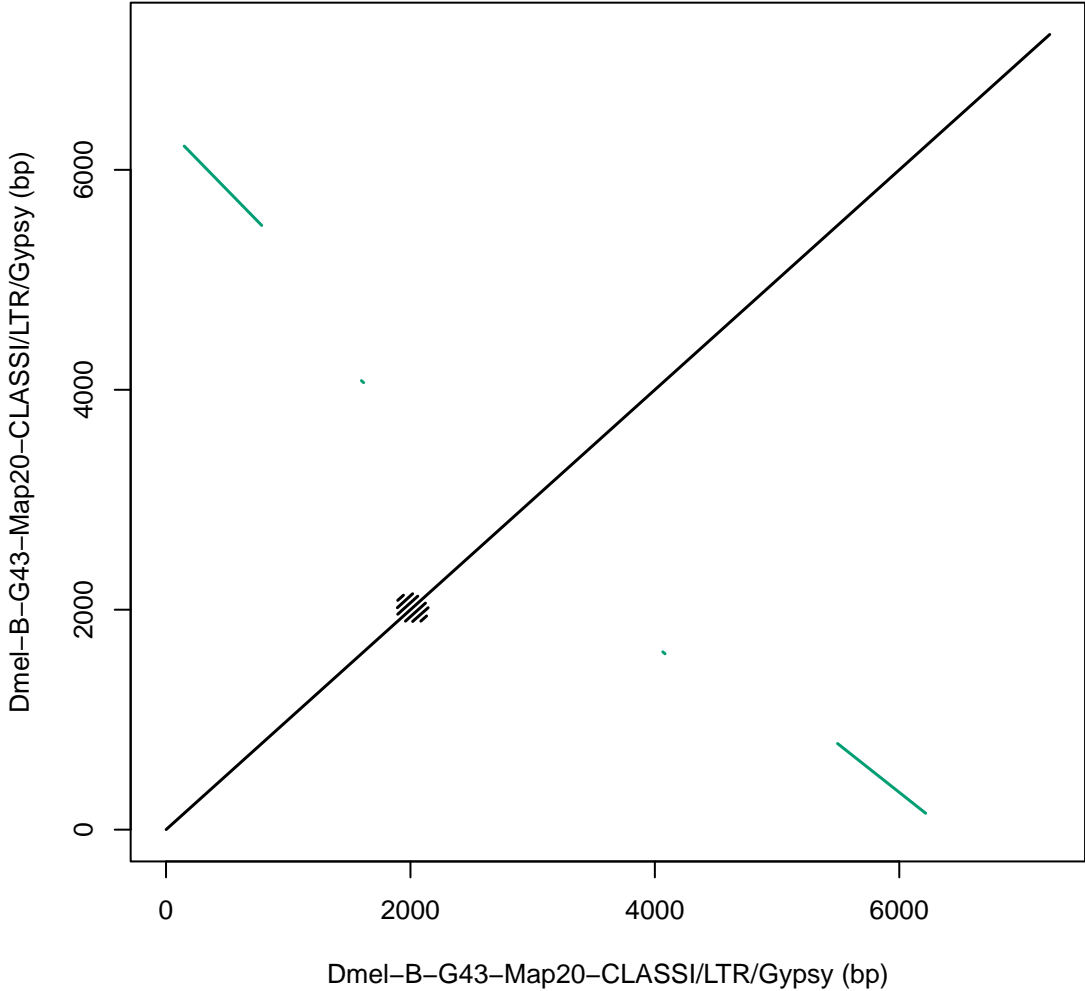
TE: Dmel-B-G43-Map20-CLASSI/LTR/Gypsy
consensus size: 7231bp; fragments: 1049; full length: 1 (>=6507.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

