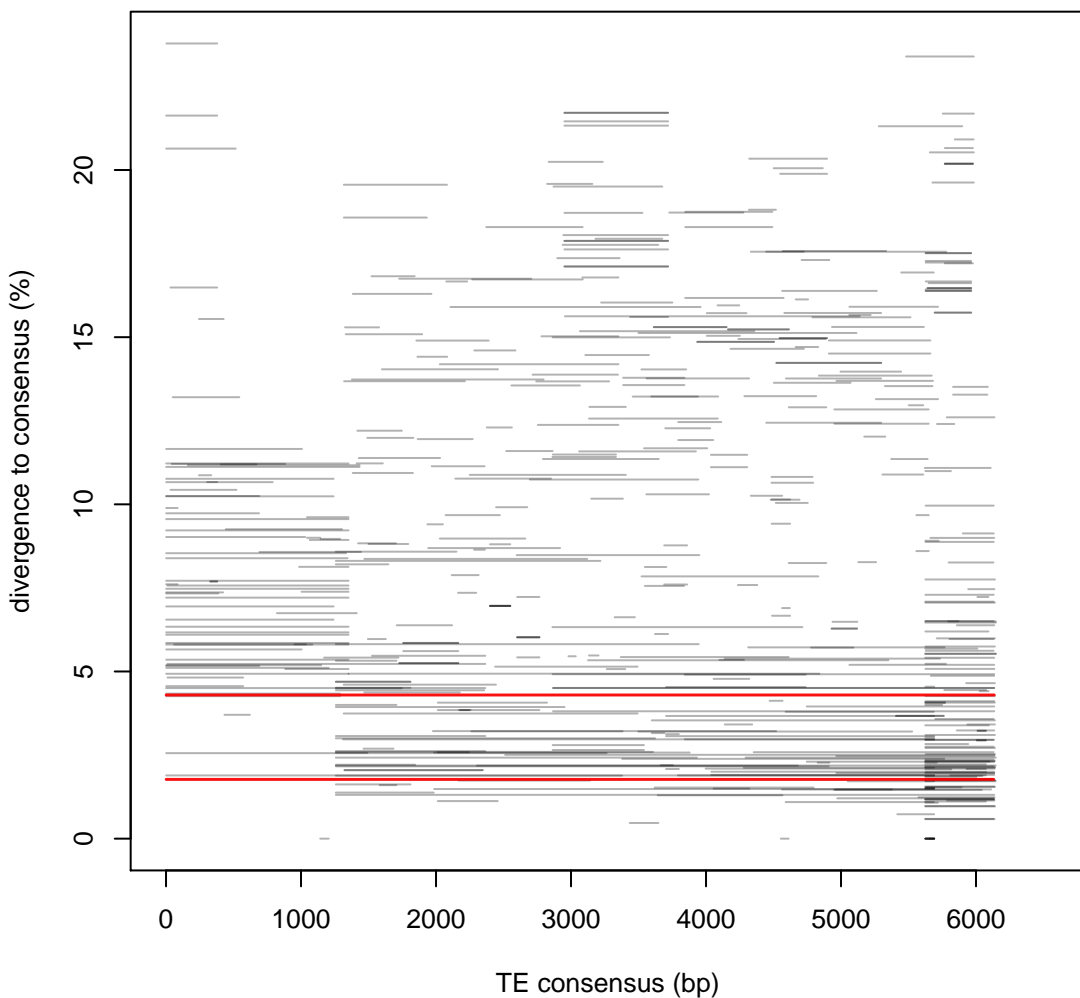
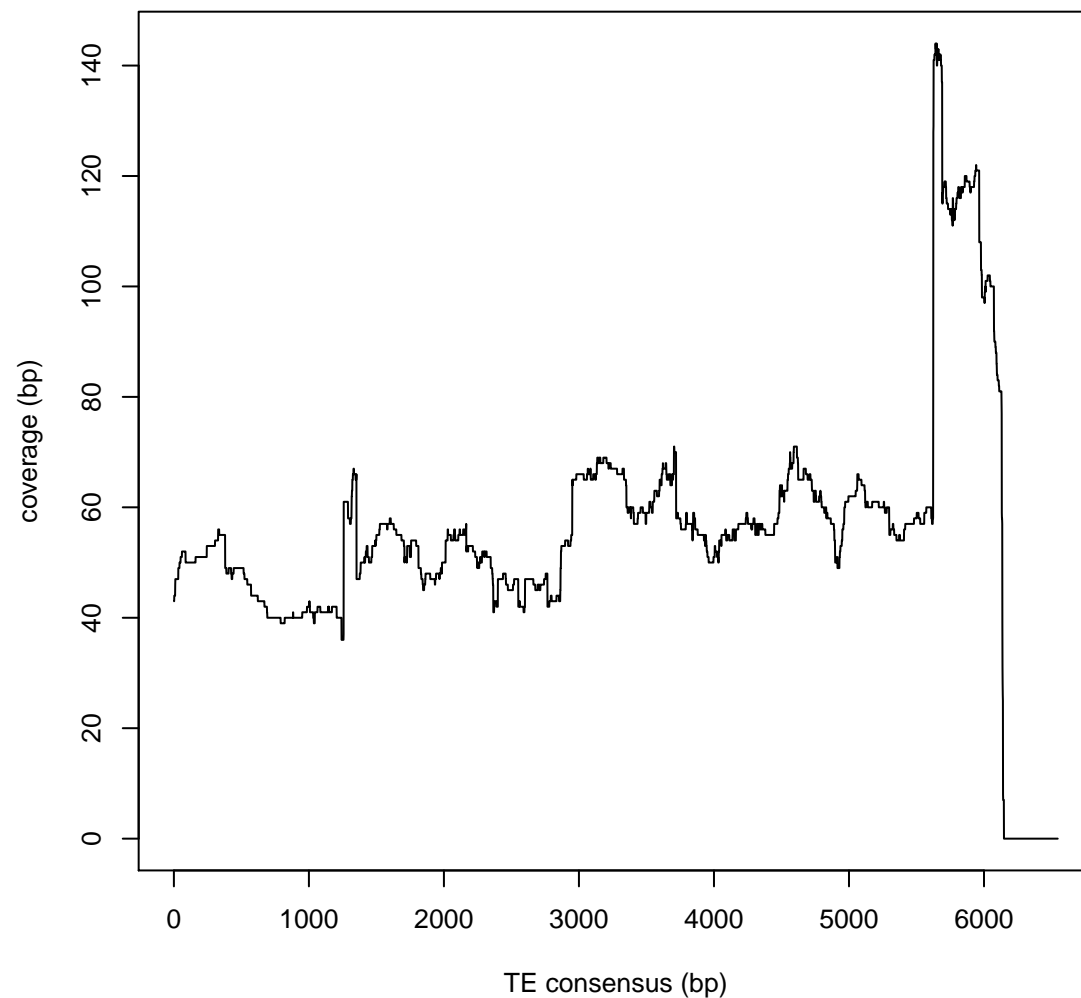


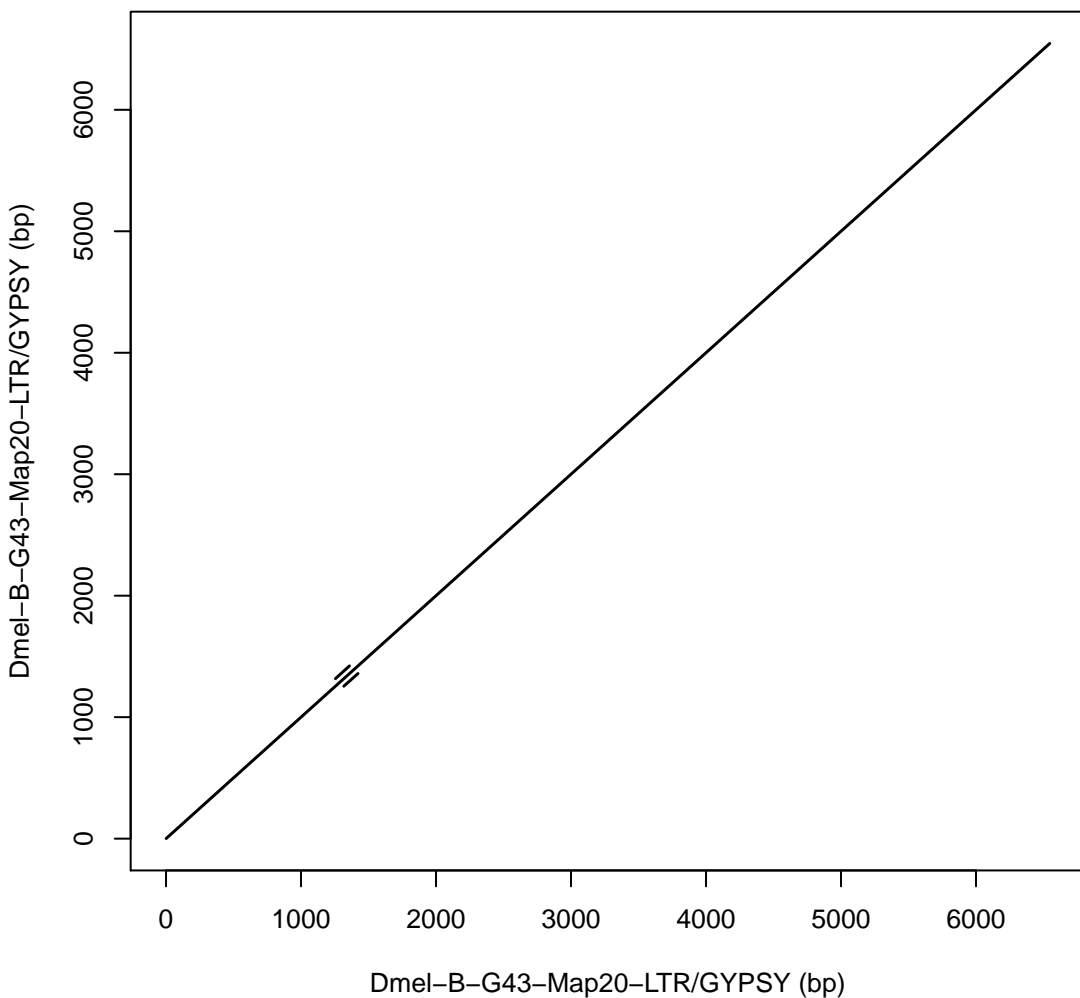
TE: Dmel-B-G43-Map20-LTR/GYPSY
consensus size: 6546bp; fragments: 498; full length: 2 (>=5891.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

