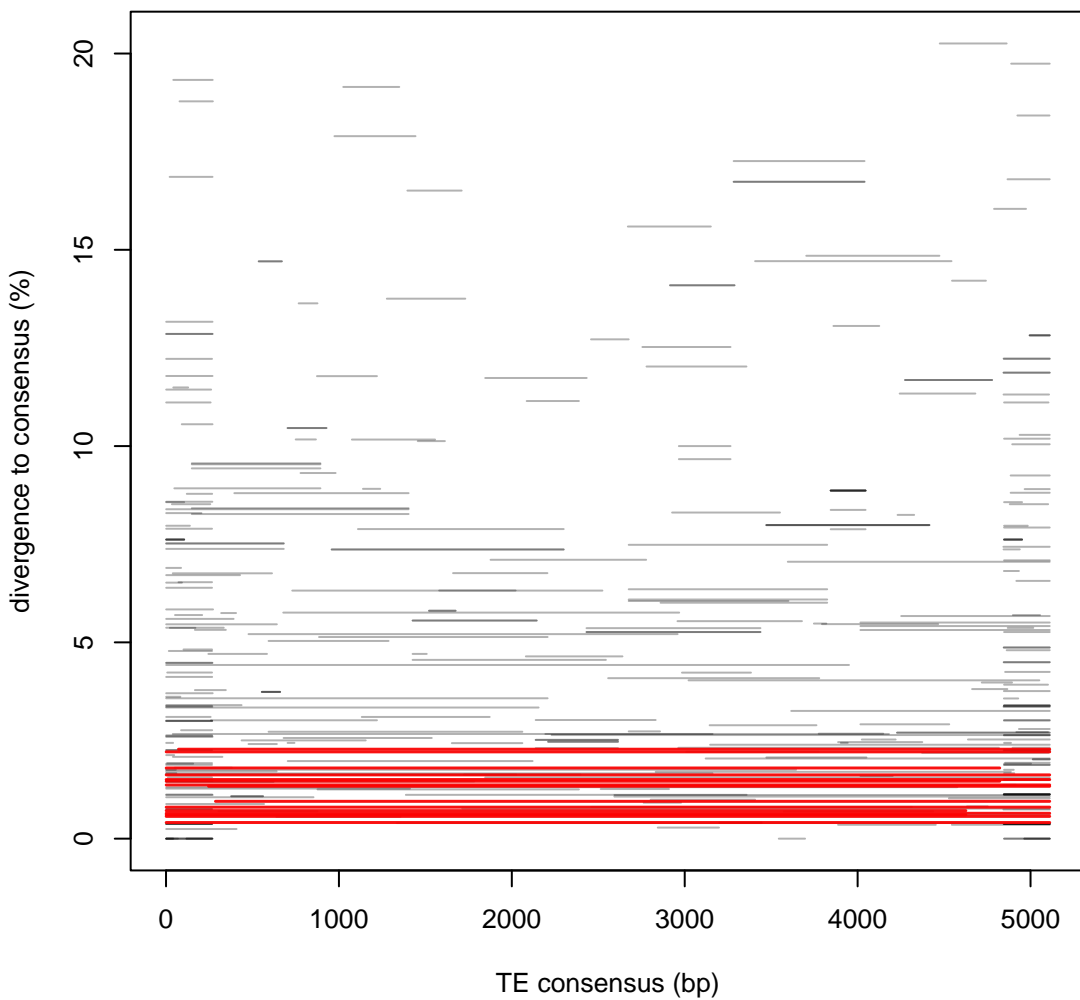
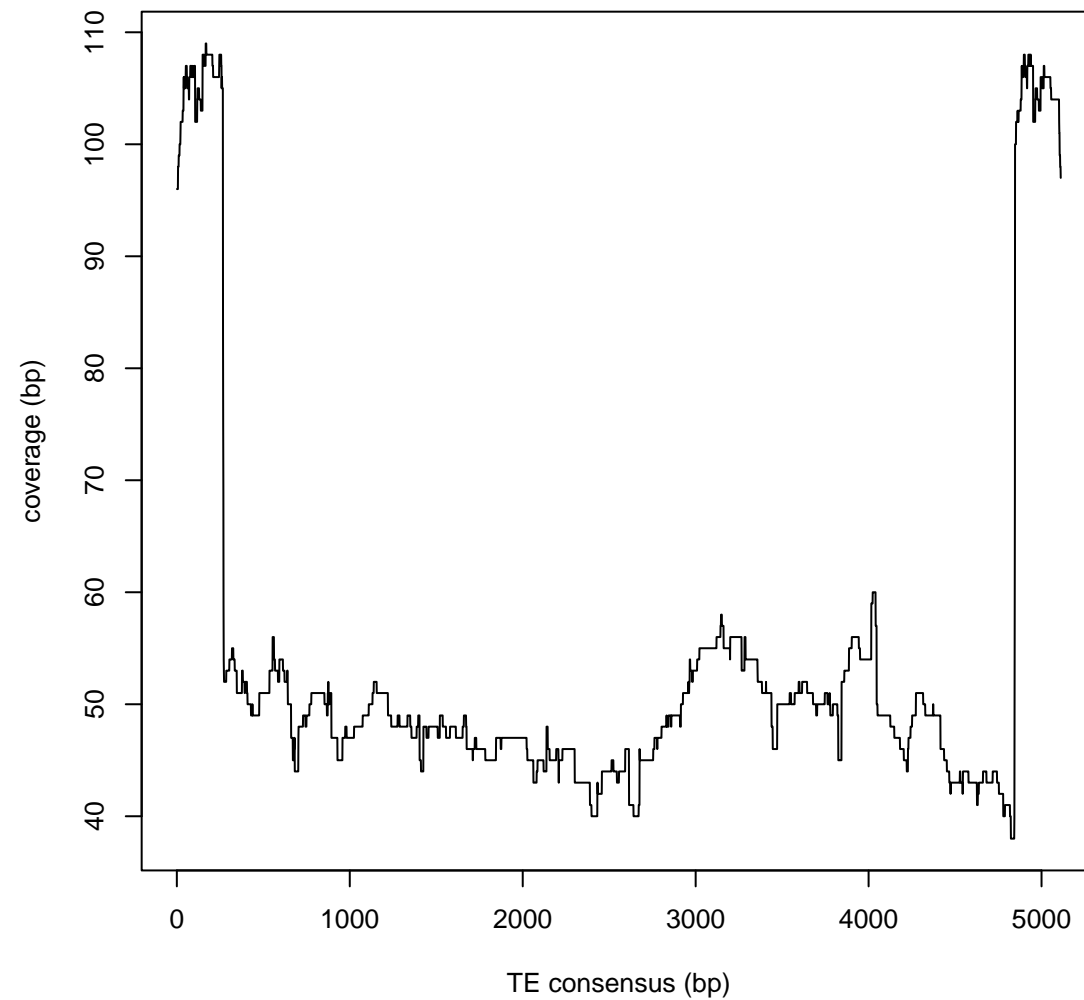


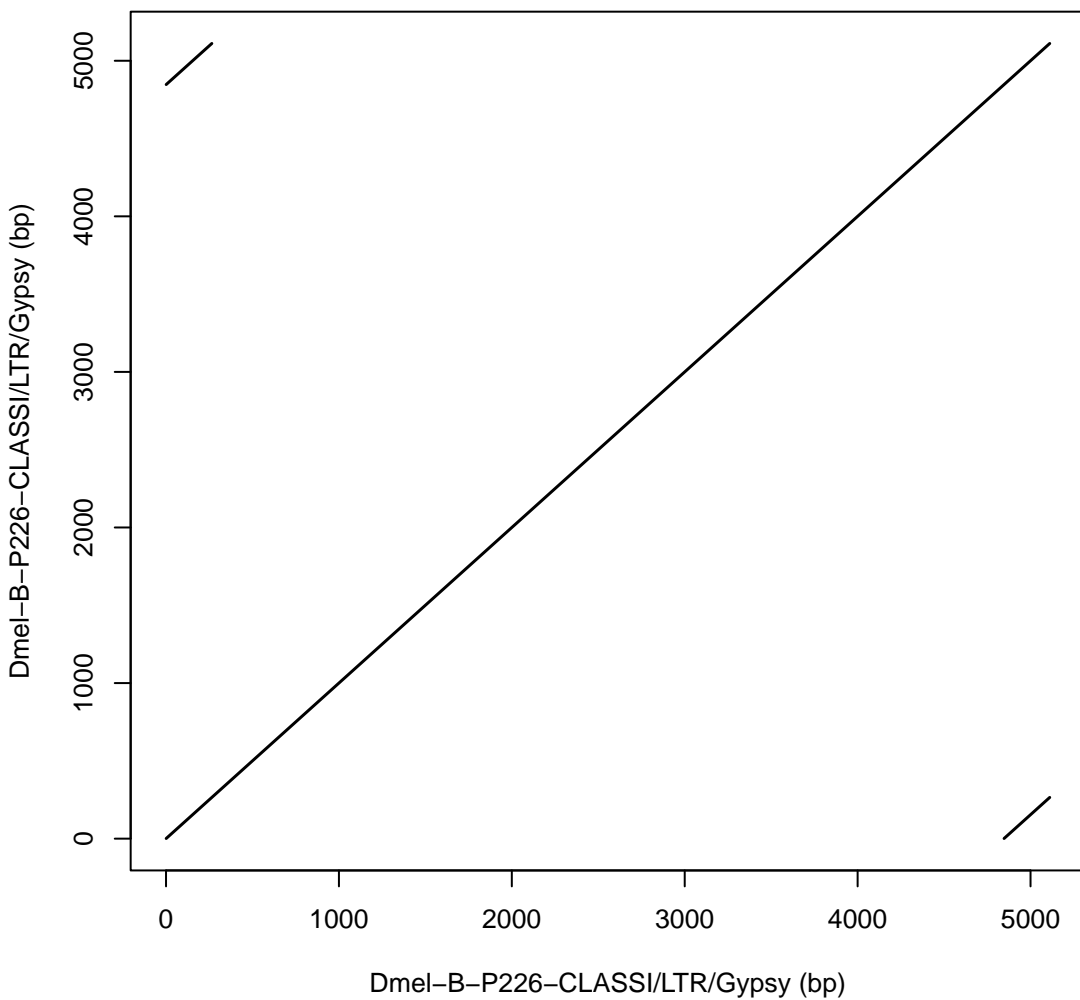
TE: Dmel-B-P226-CLASSI/LTR/Gypsy
consensus size: 5111bp; fragments: 373; full length: 15 (>=4599.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

