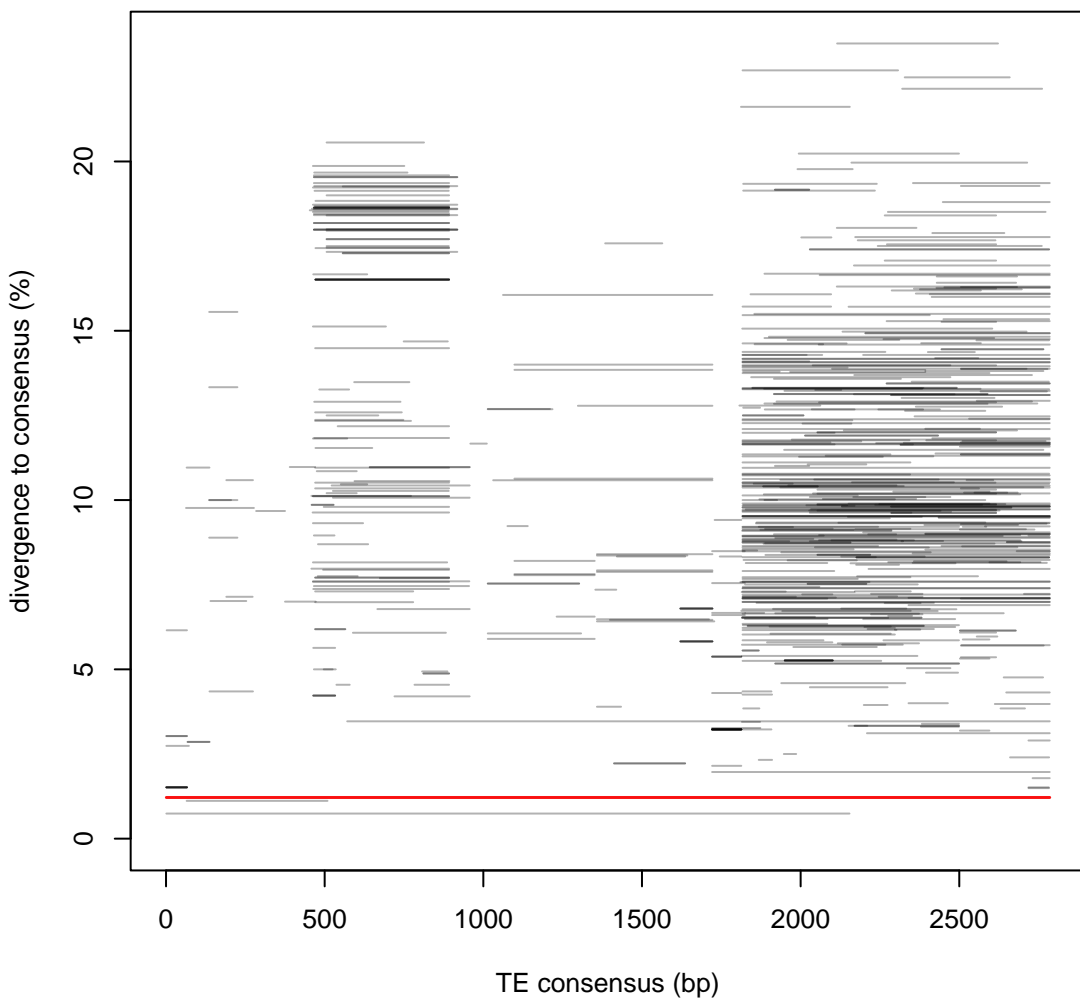
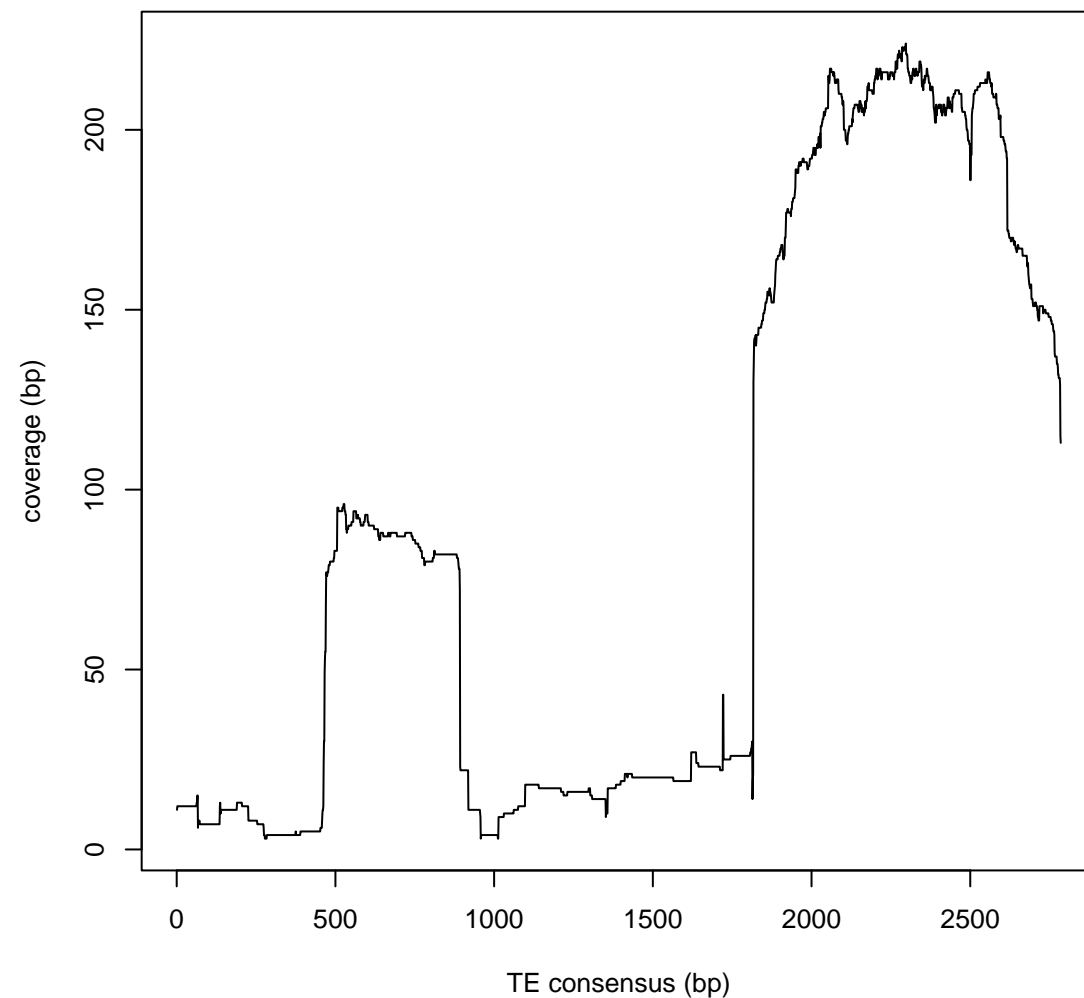


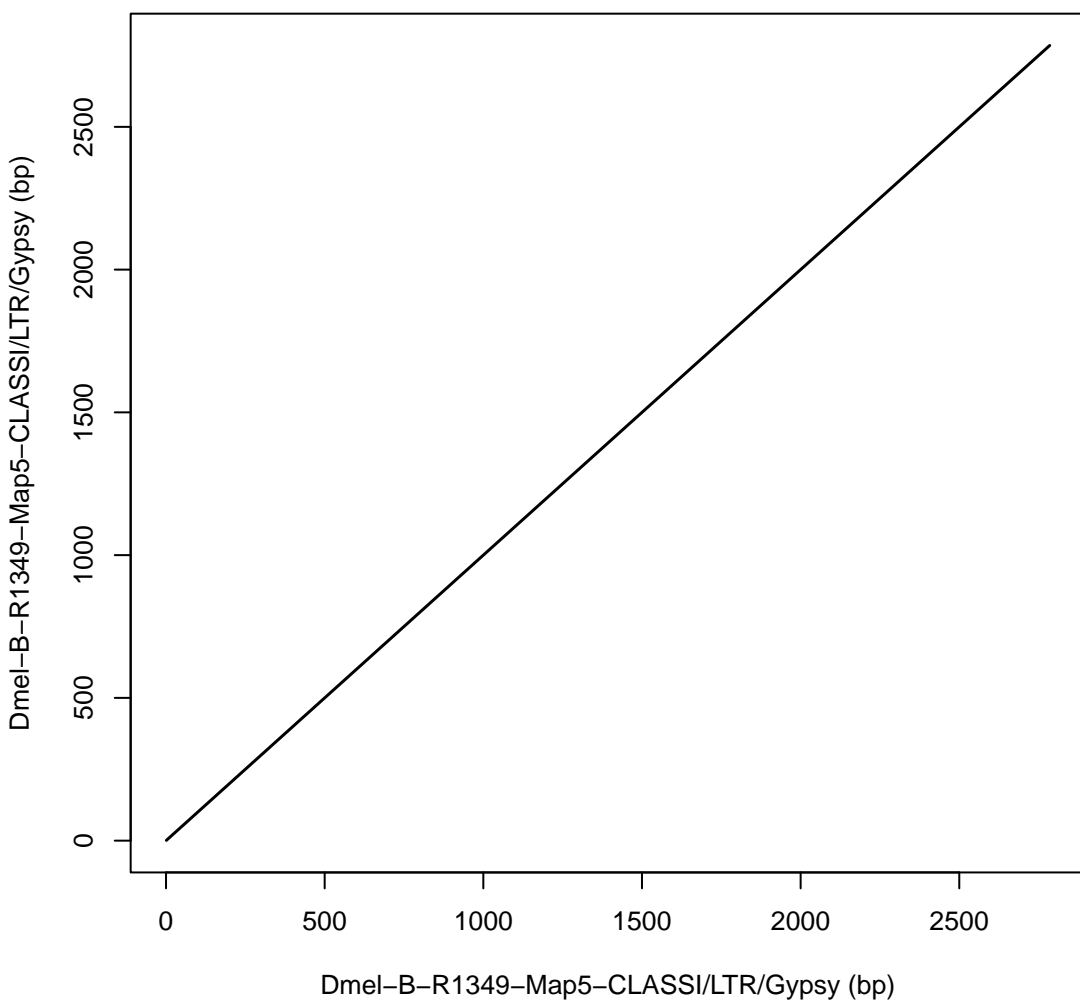
TE: Dmel-B-R1349-Map5-CLASSI/LTR/Gypsy
consensus size: 2785bp; fragments: 629; full length: 1 (>=2506.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

