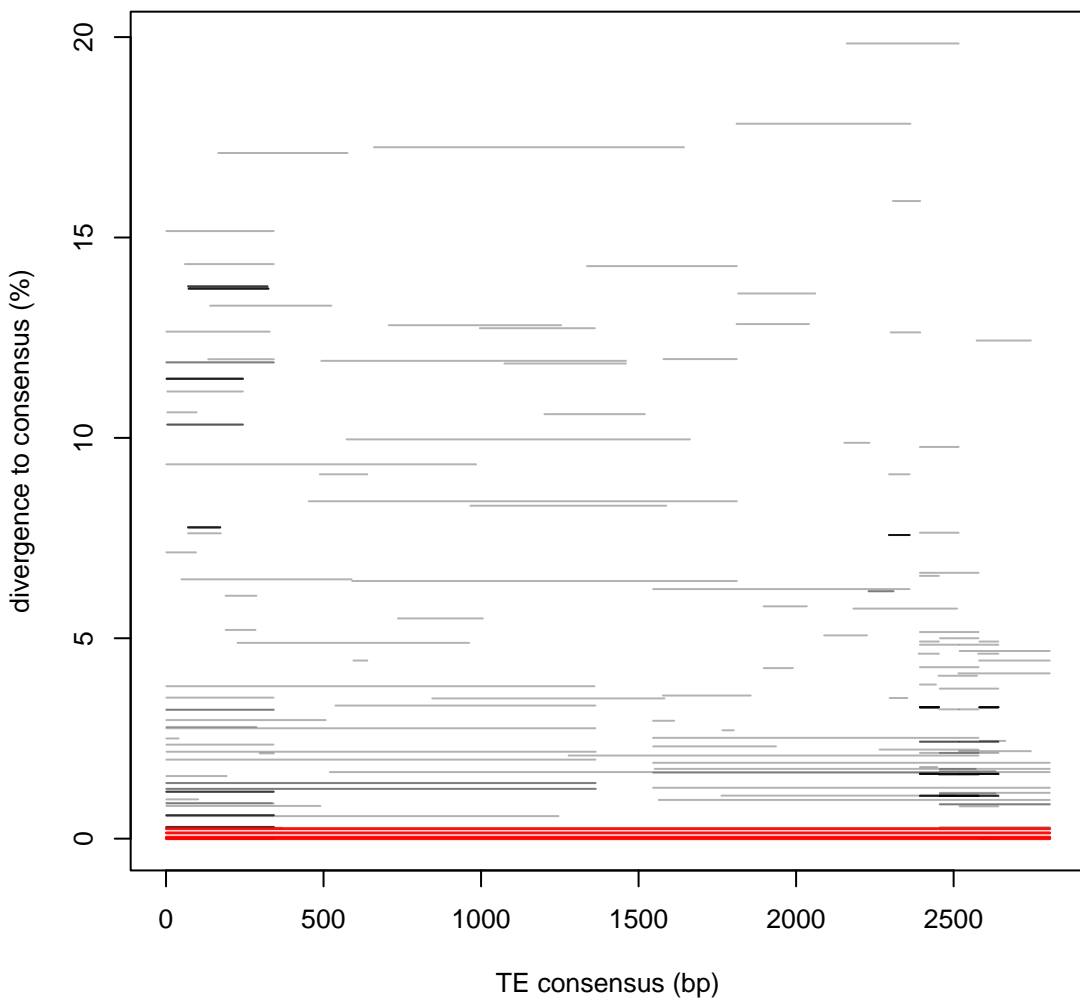
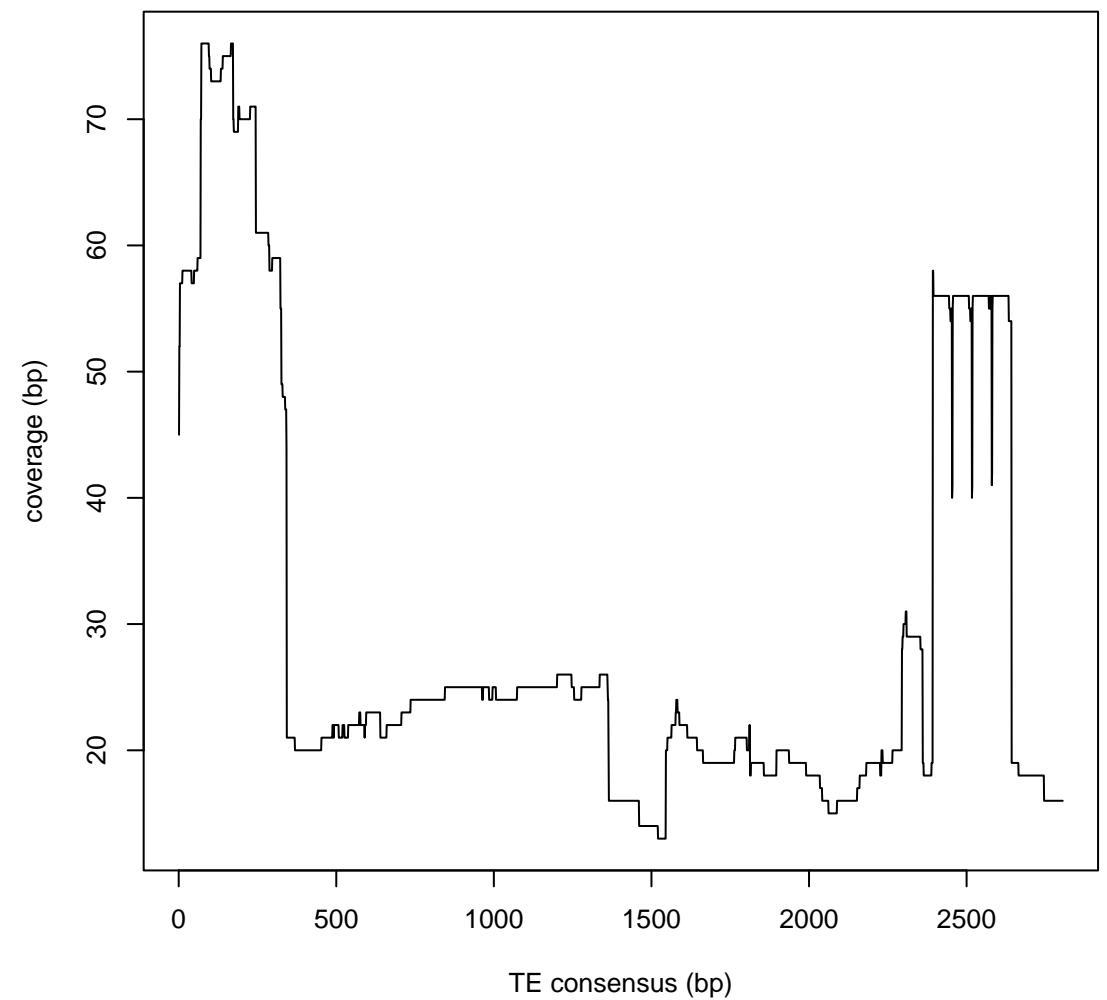


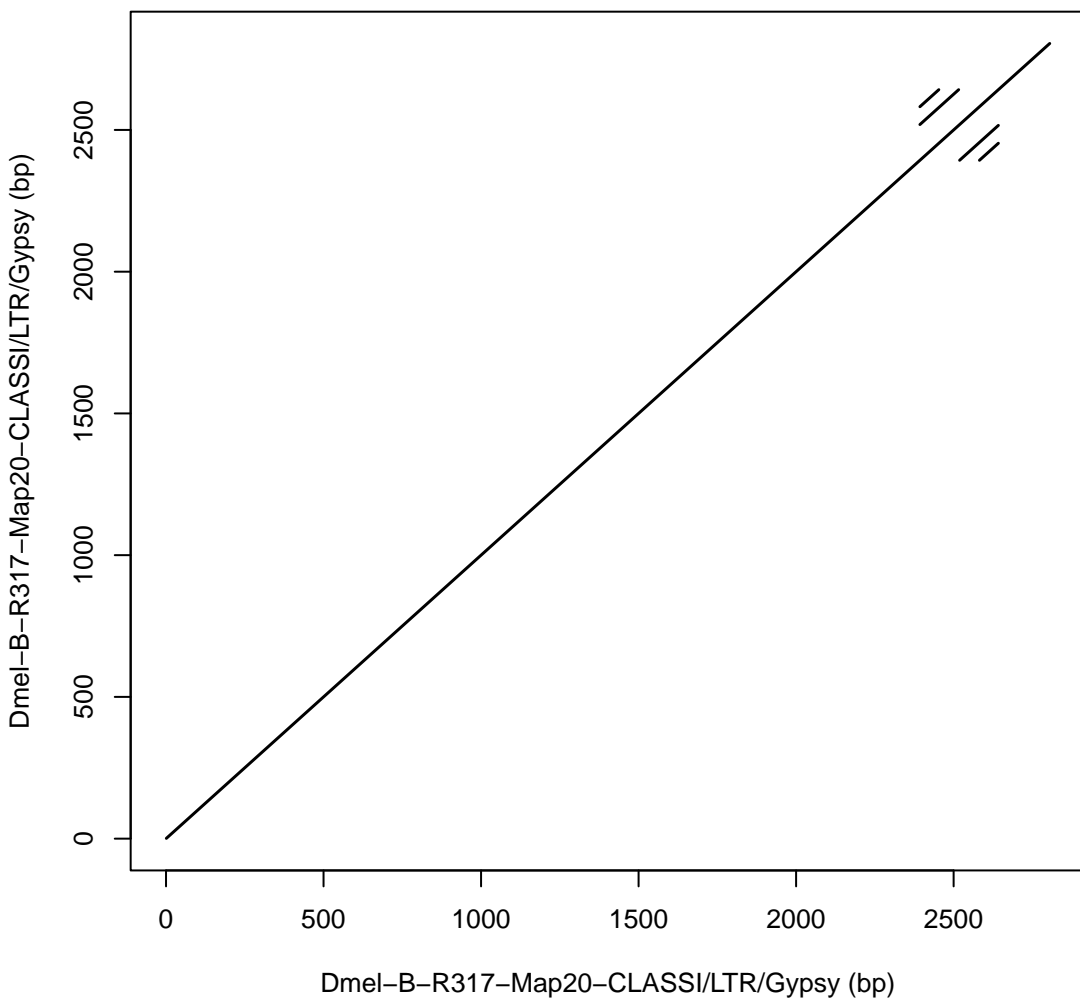
TE: Dmel-B-R317-Map20-CLASSI/LTR/Gypsy
consensus size: 2805bp; fragments: 232; full length: 4 (>=2524.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

