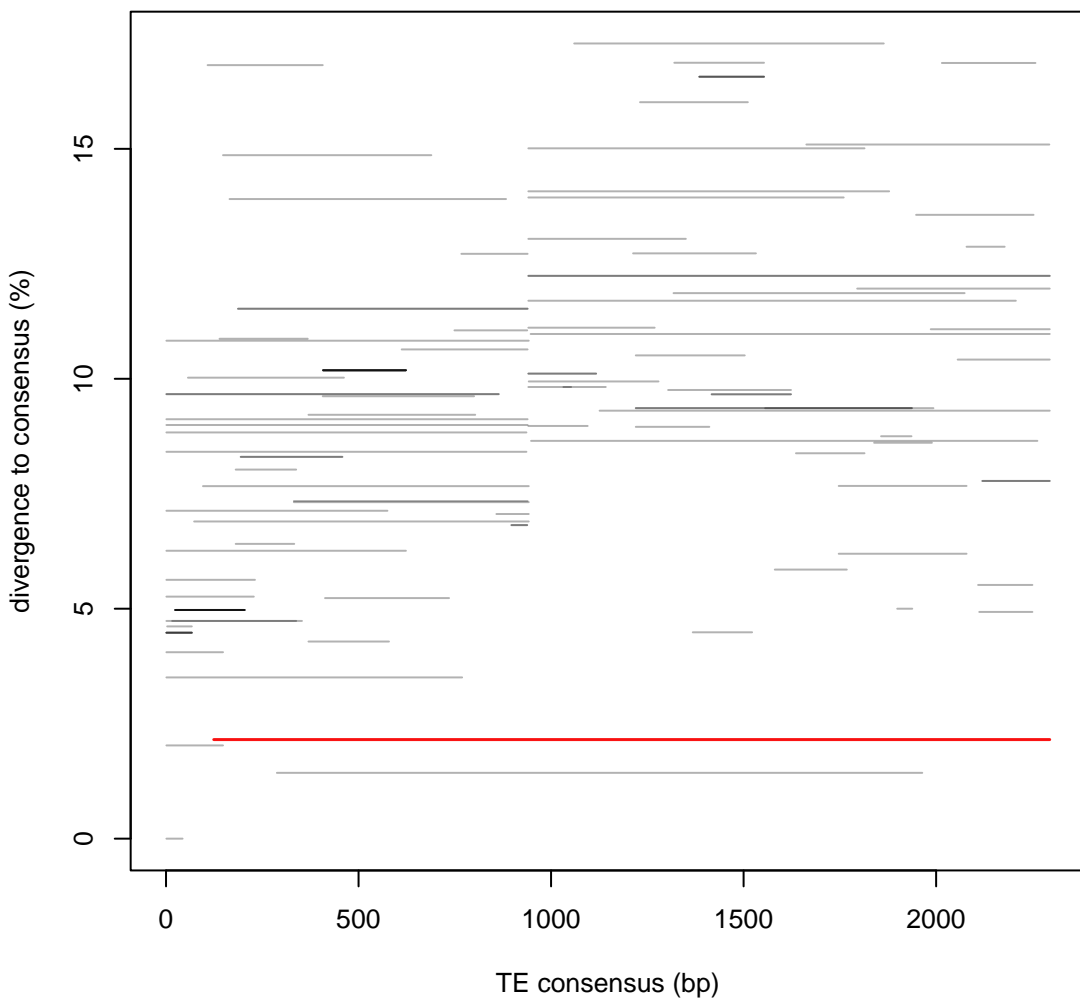
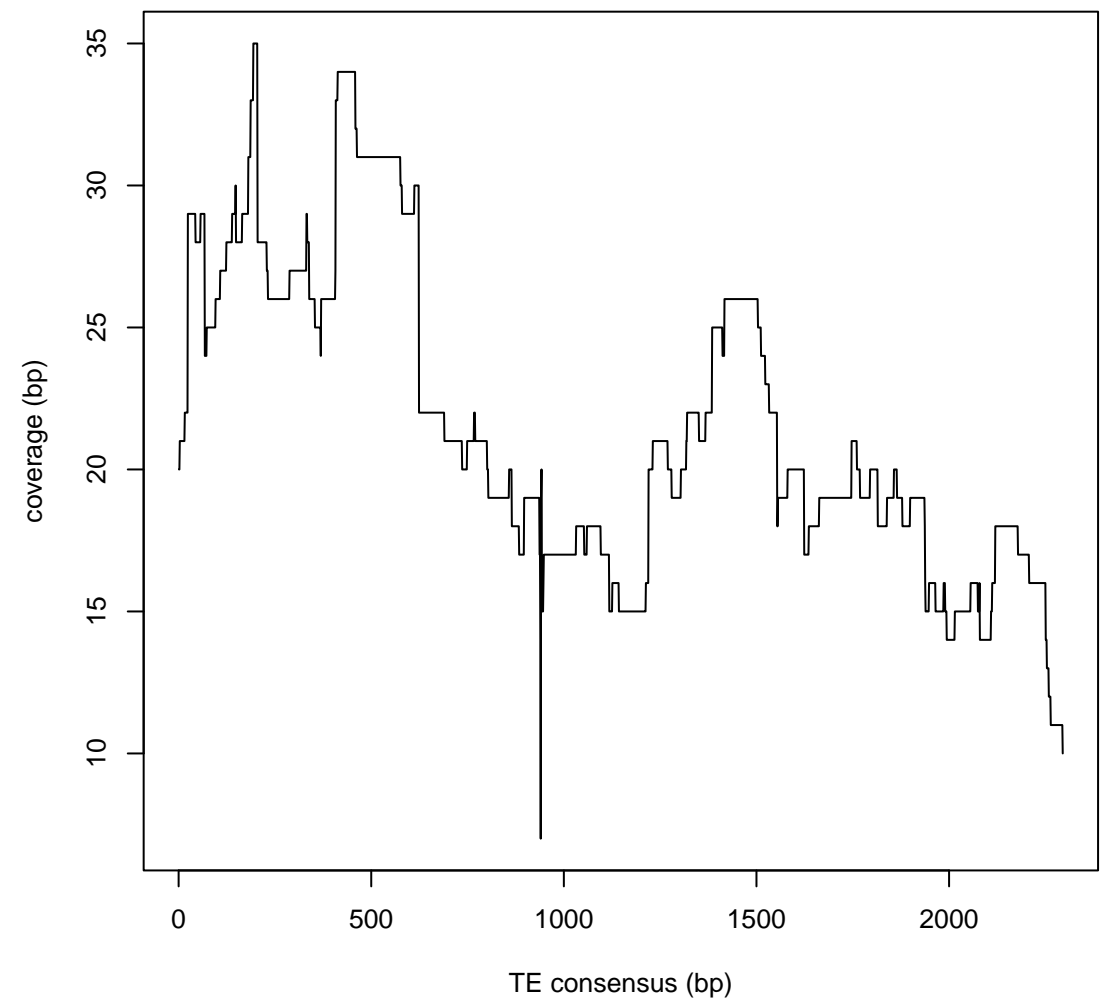


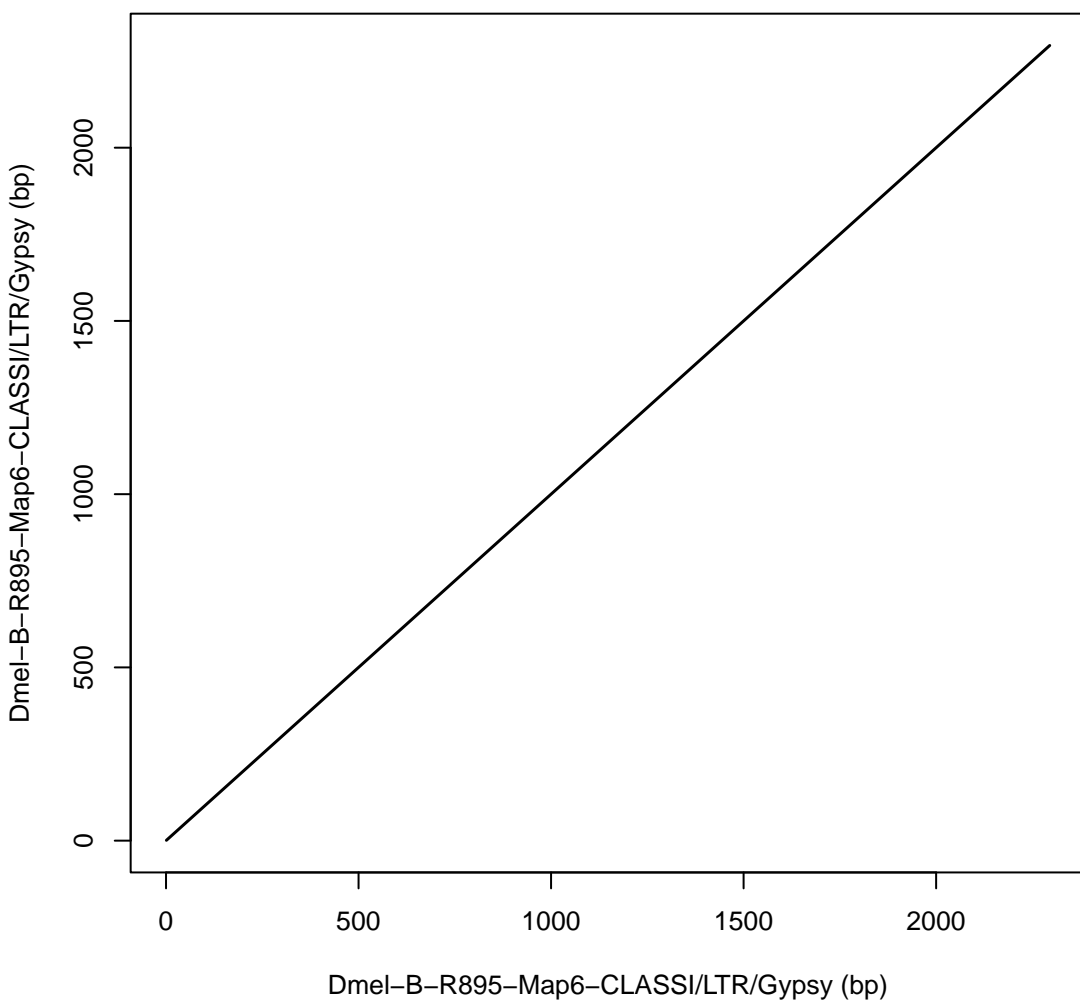
TE: Dmel-B-R895-Map6-CLASSI/LTR/Gypsy
consensus size: 2295bp; fragments: 115; full length: 1 (>=2065.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

