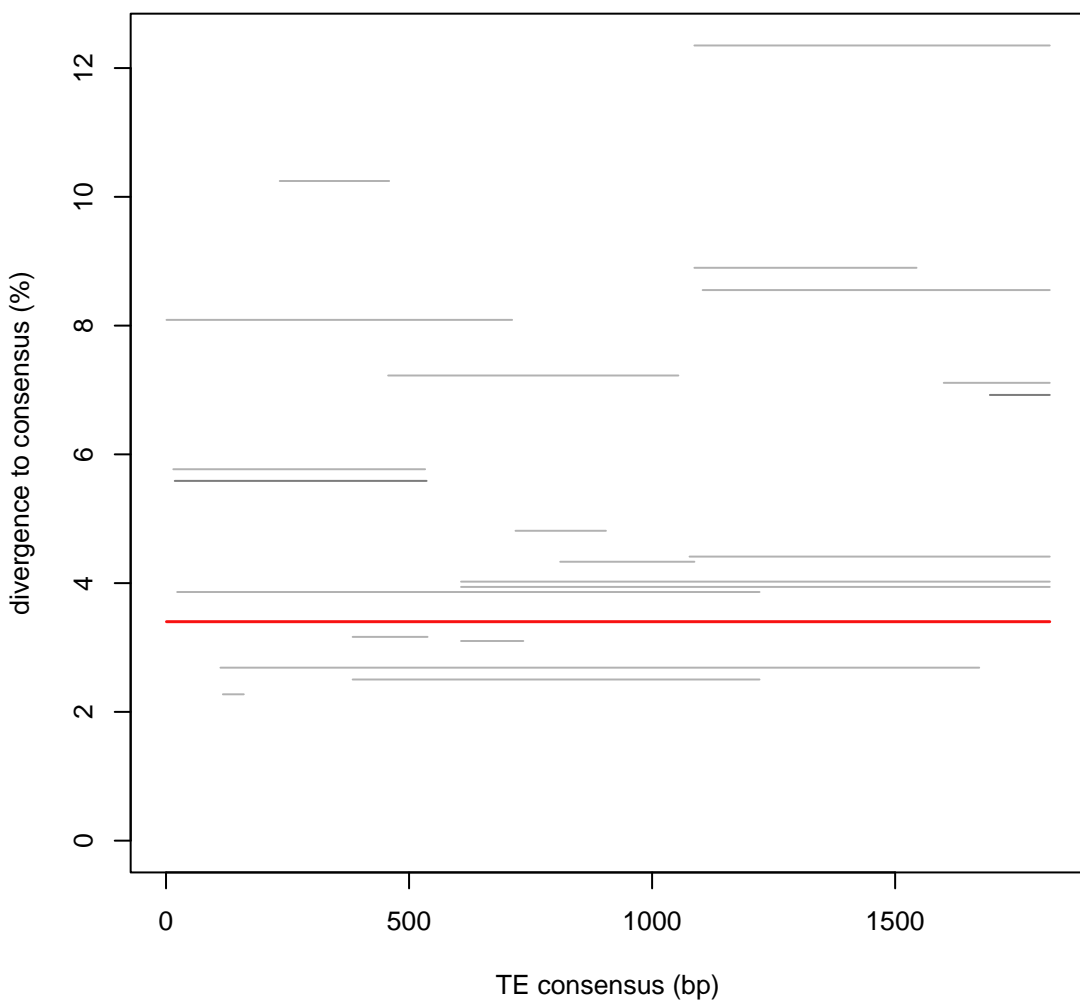
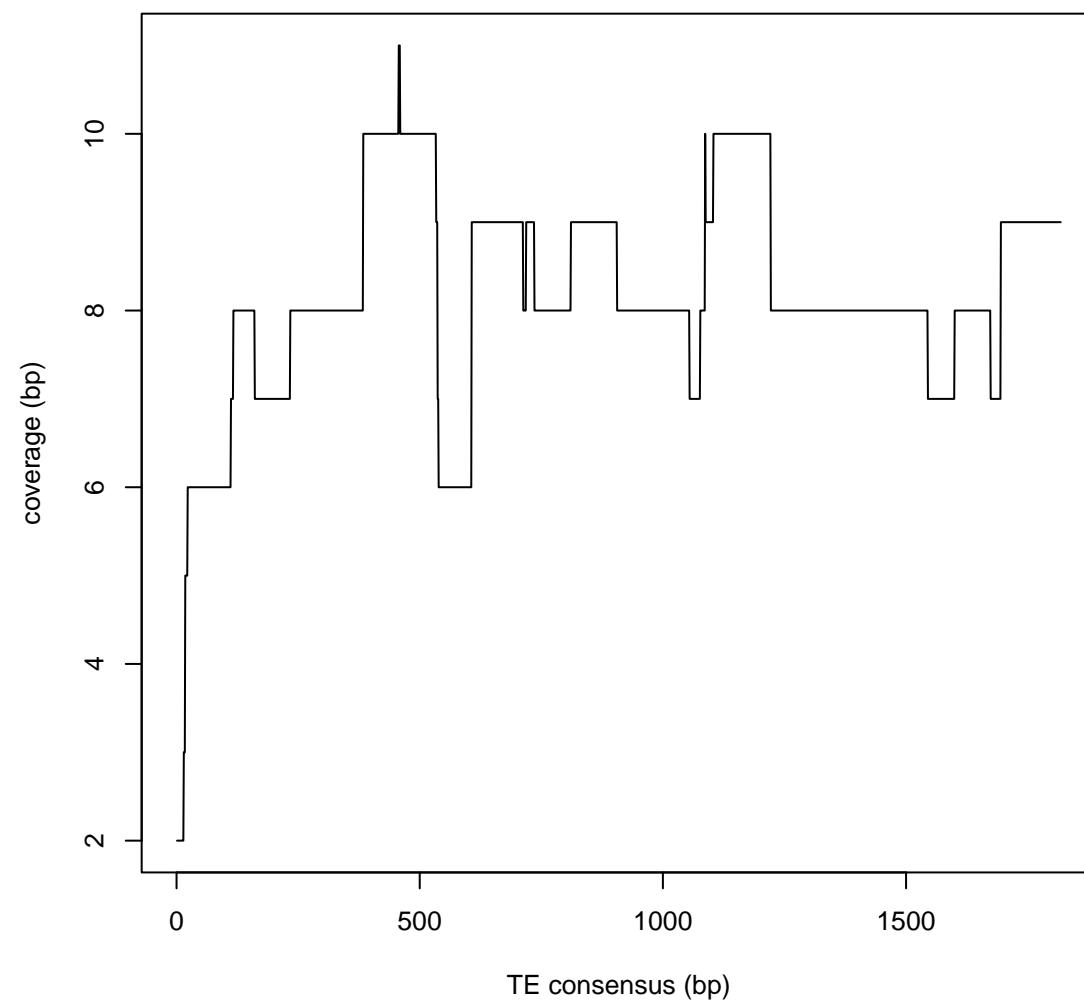


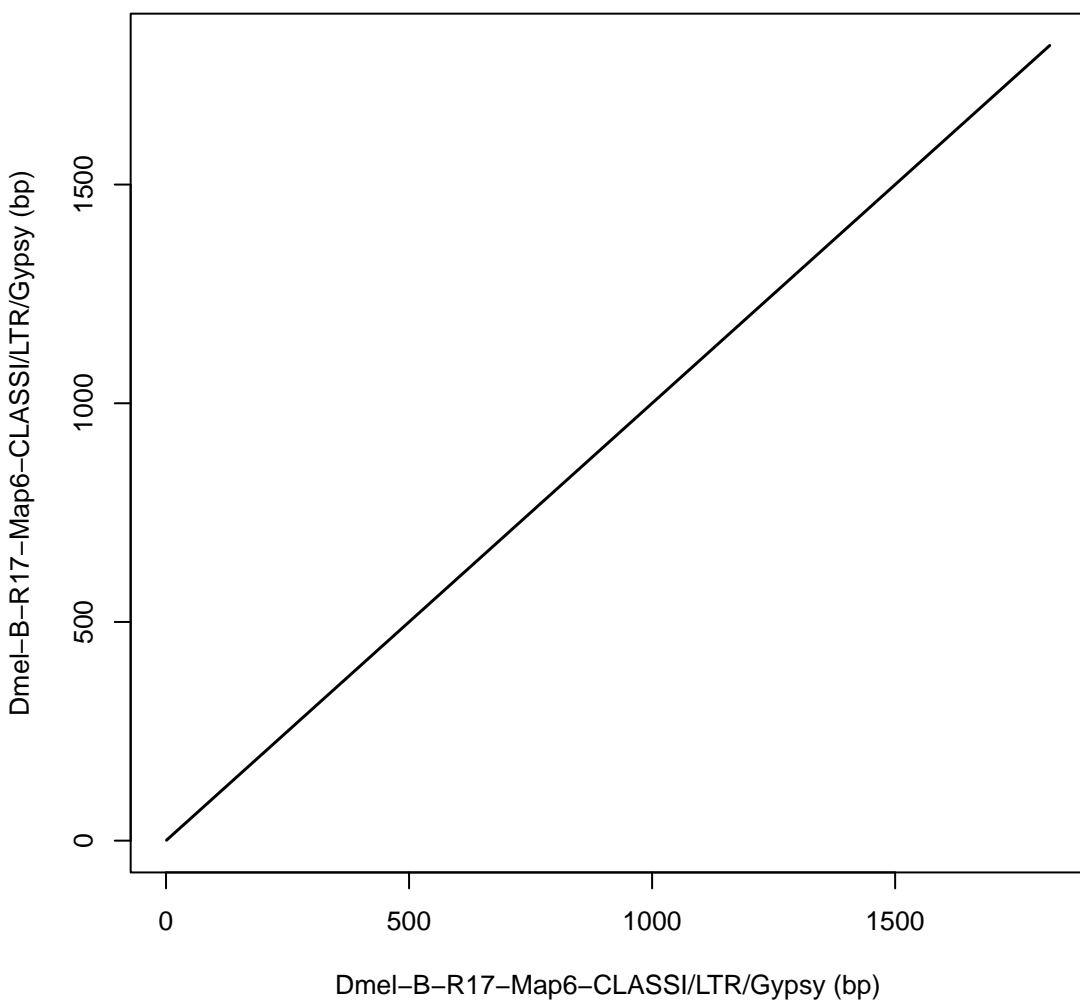
TE: Dmel-B-R17-Map6-CLASSI/LTR/Gypsy
consensus size: 1818bp; fragments: 24; full length: 1 (>=1636.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

