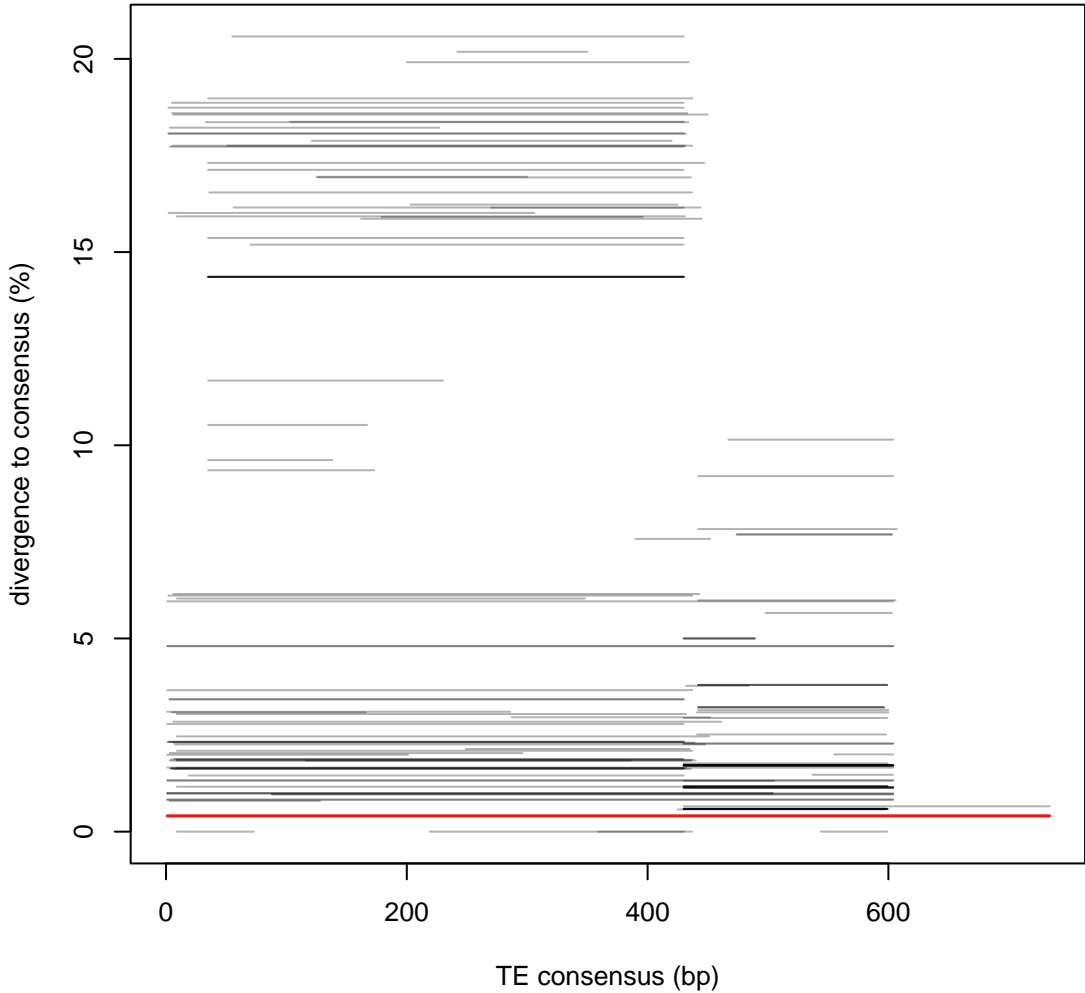
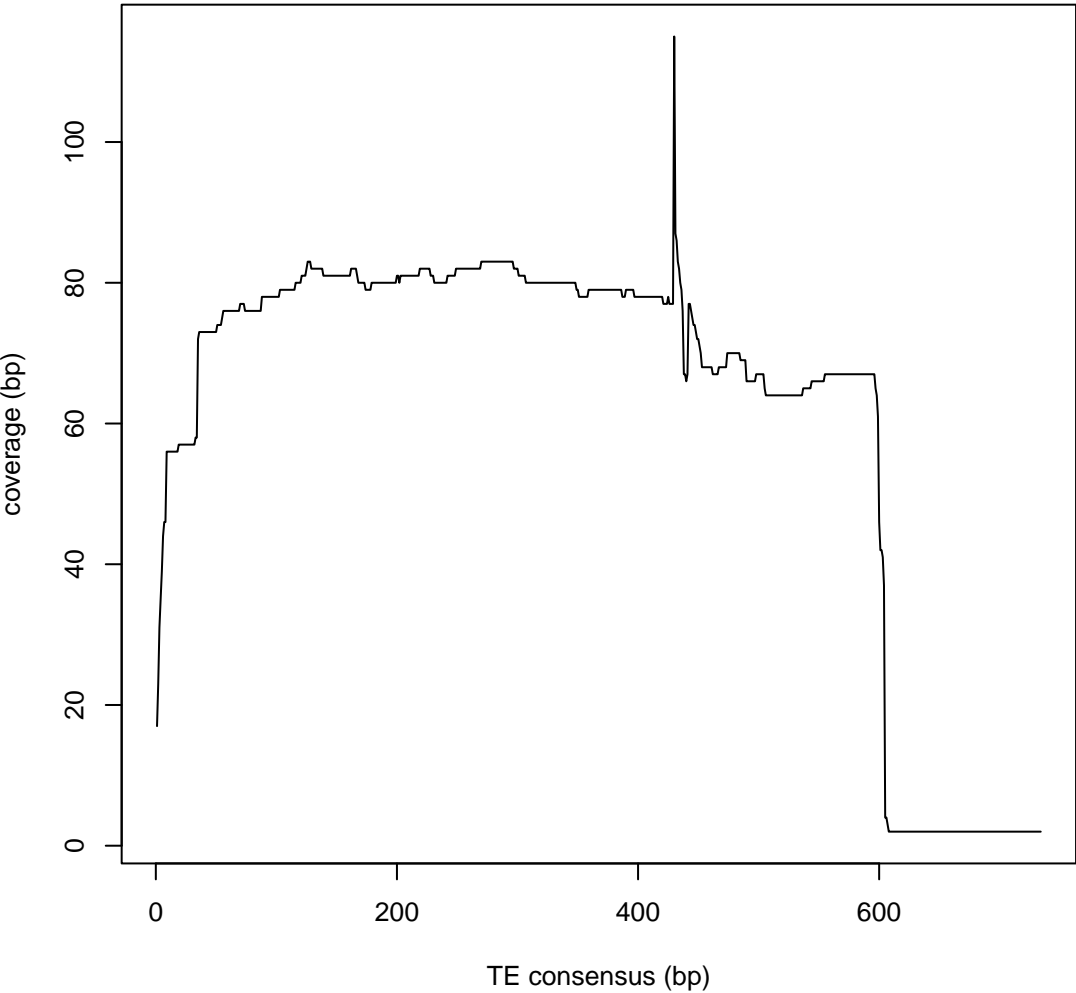


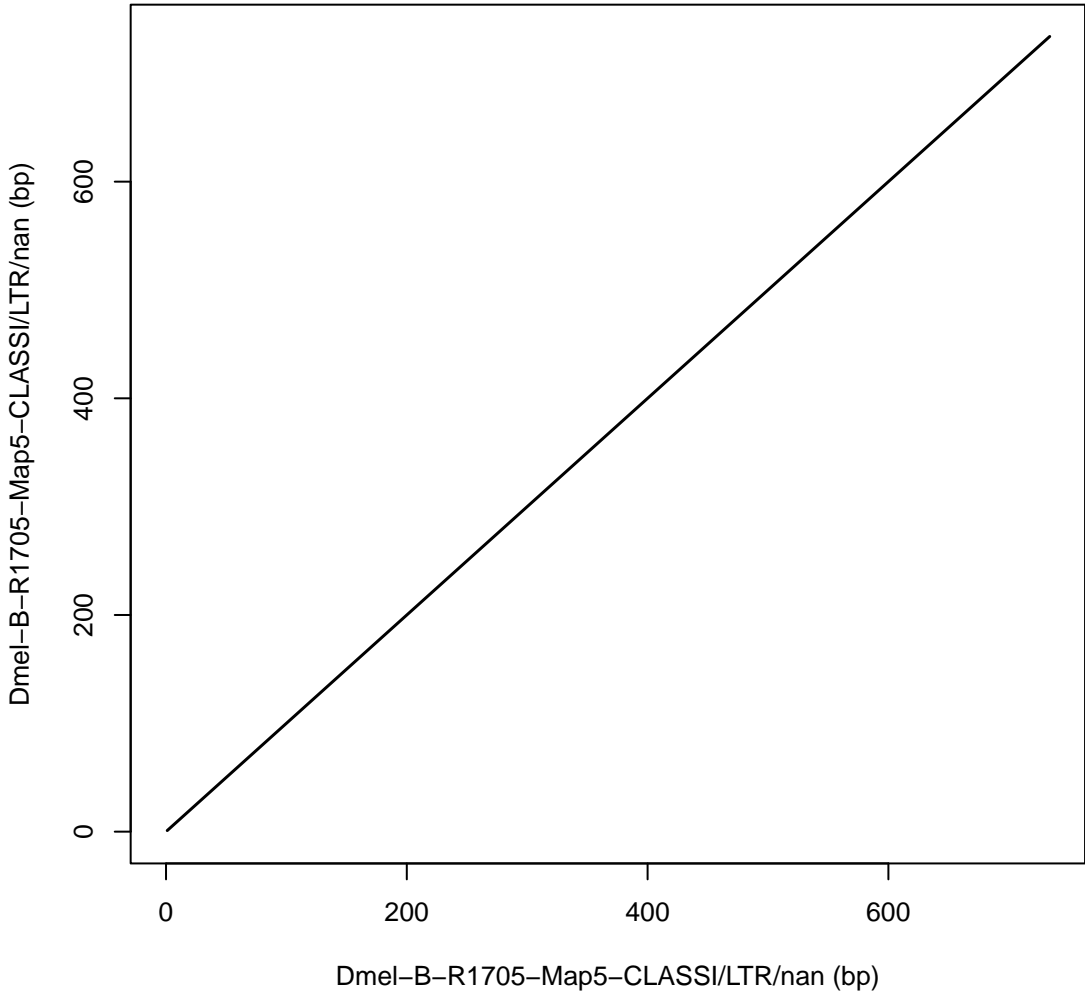
TE: Dmel-B-R1705-Map5-CLASSI/LTR/nan
consensus size: 734bp; fragments: 155; full length: 1 (>=660.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

