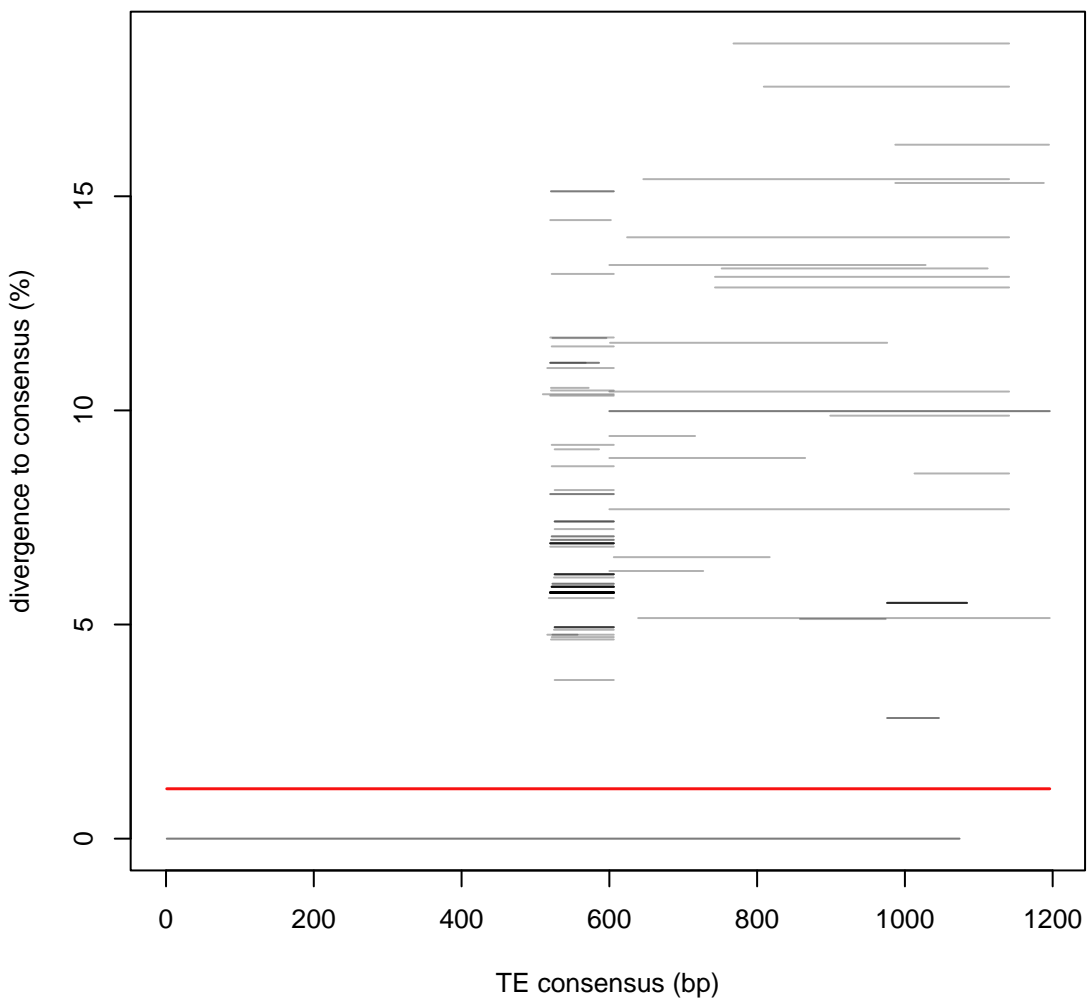
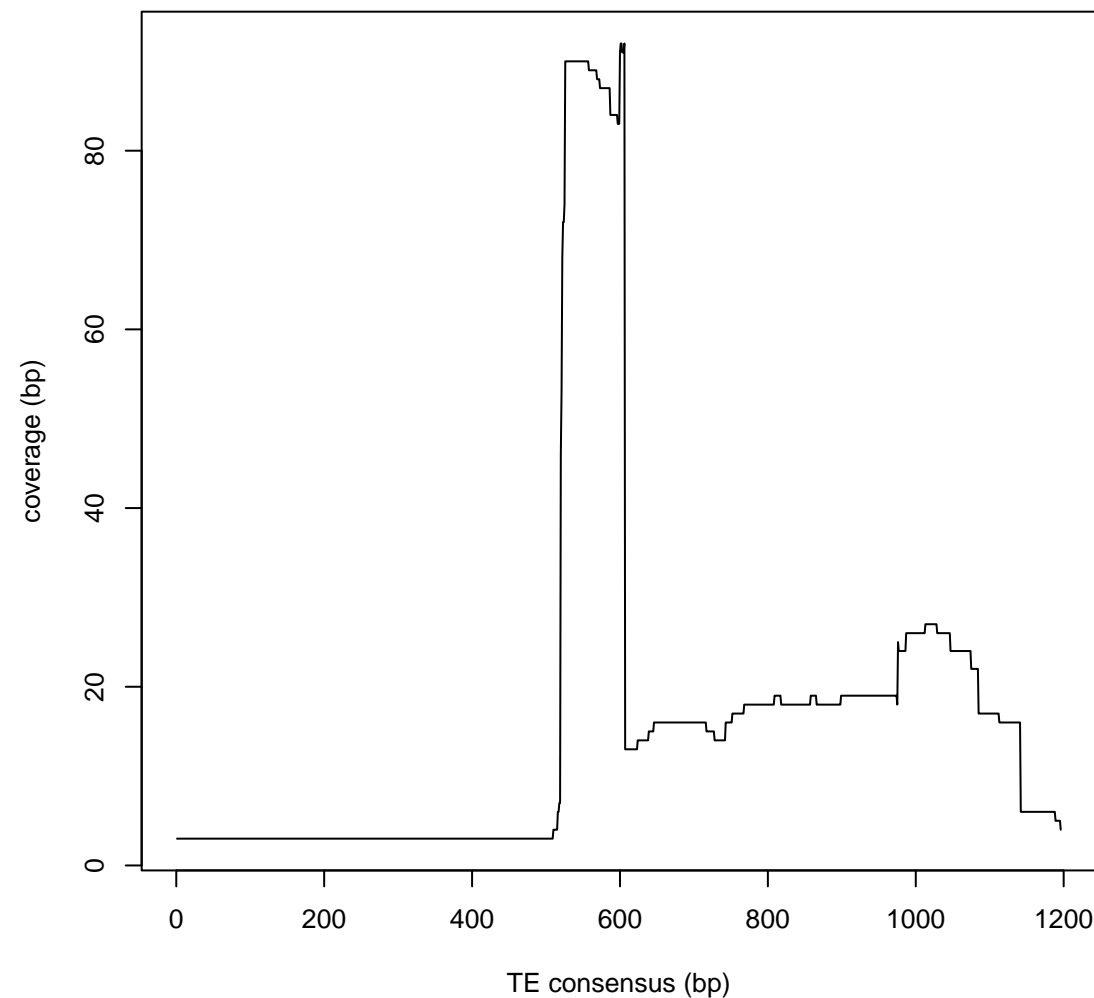


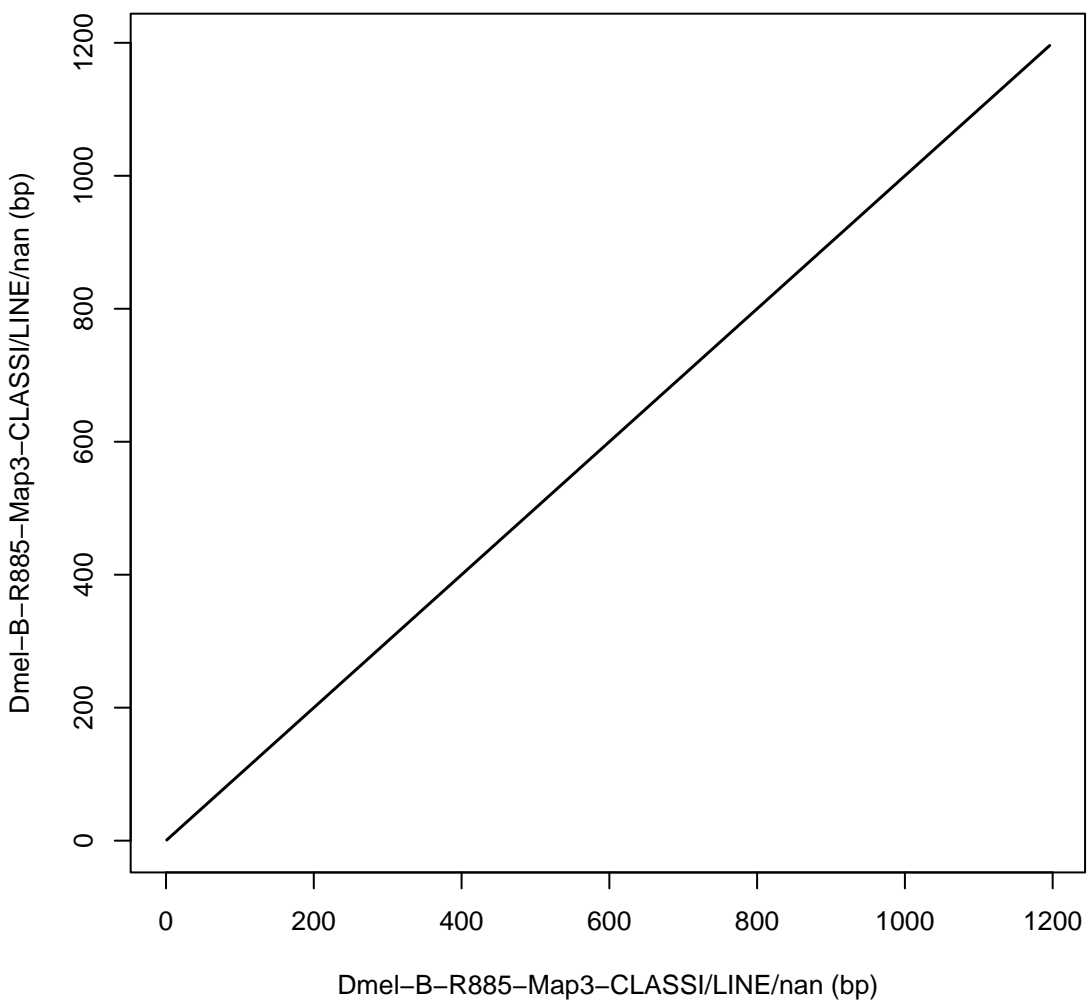
TE: Dmel-B-R885-Map3-CLASSI/LINE/nan  
consensus size: 1196bp; fragments: 120; full length: 1 ( $\geq 1076.4$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

