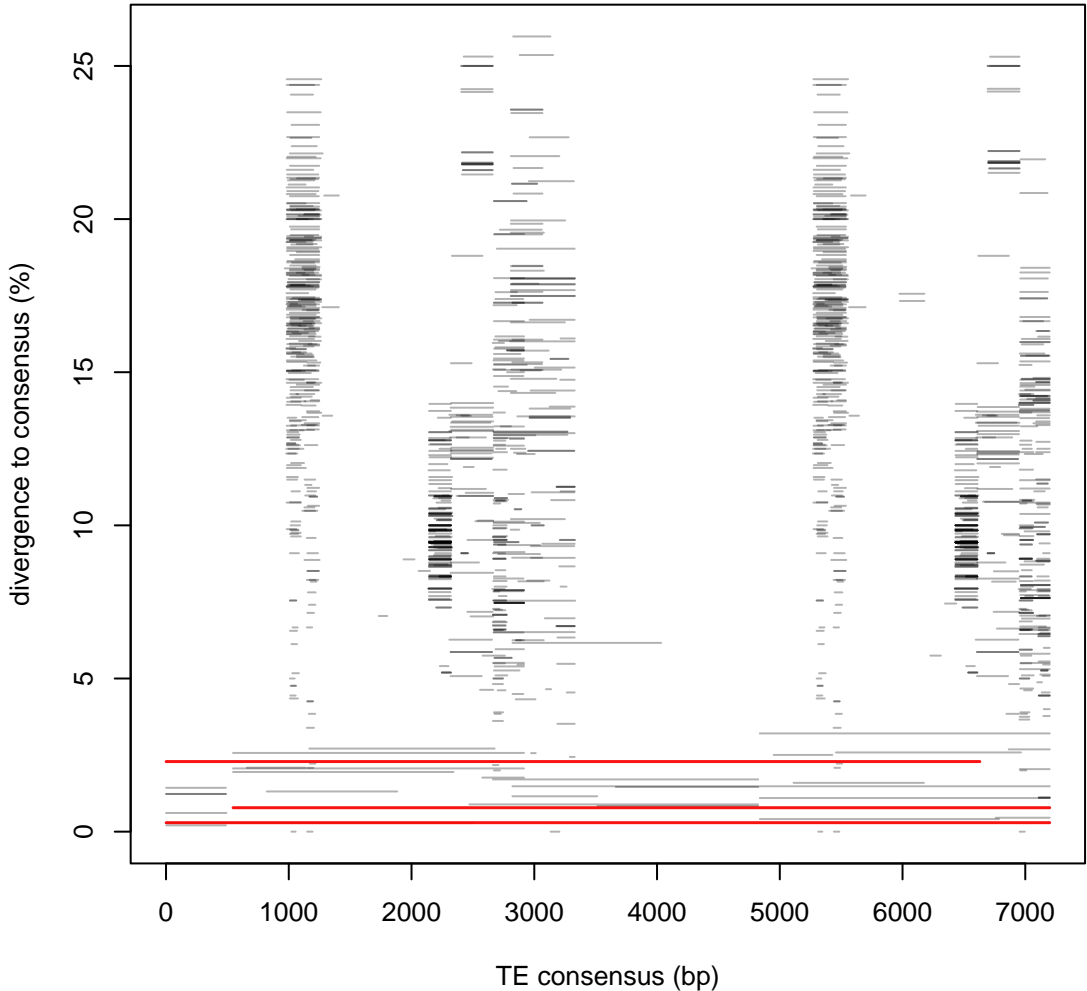
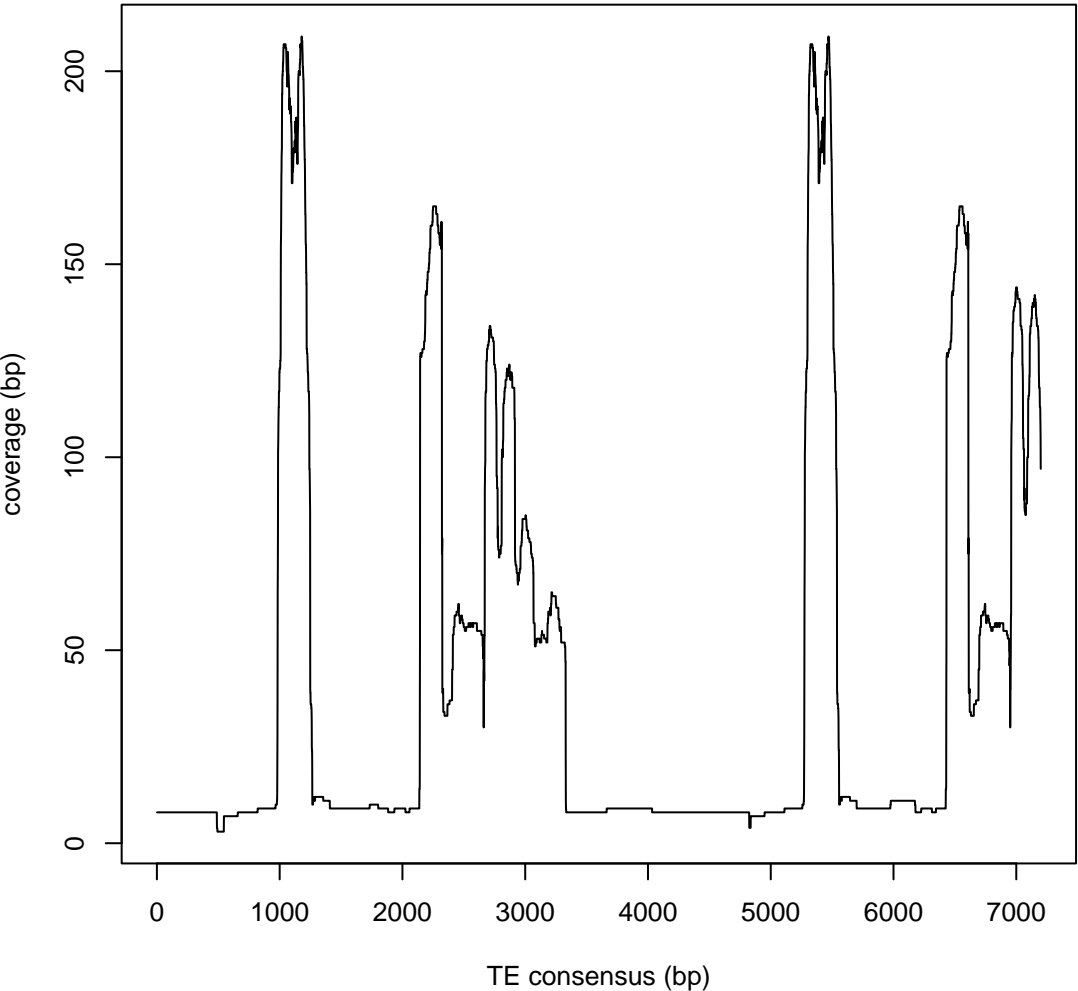


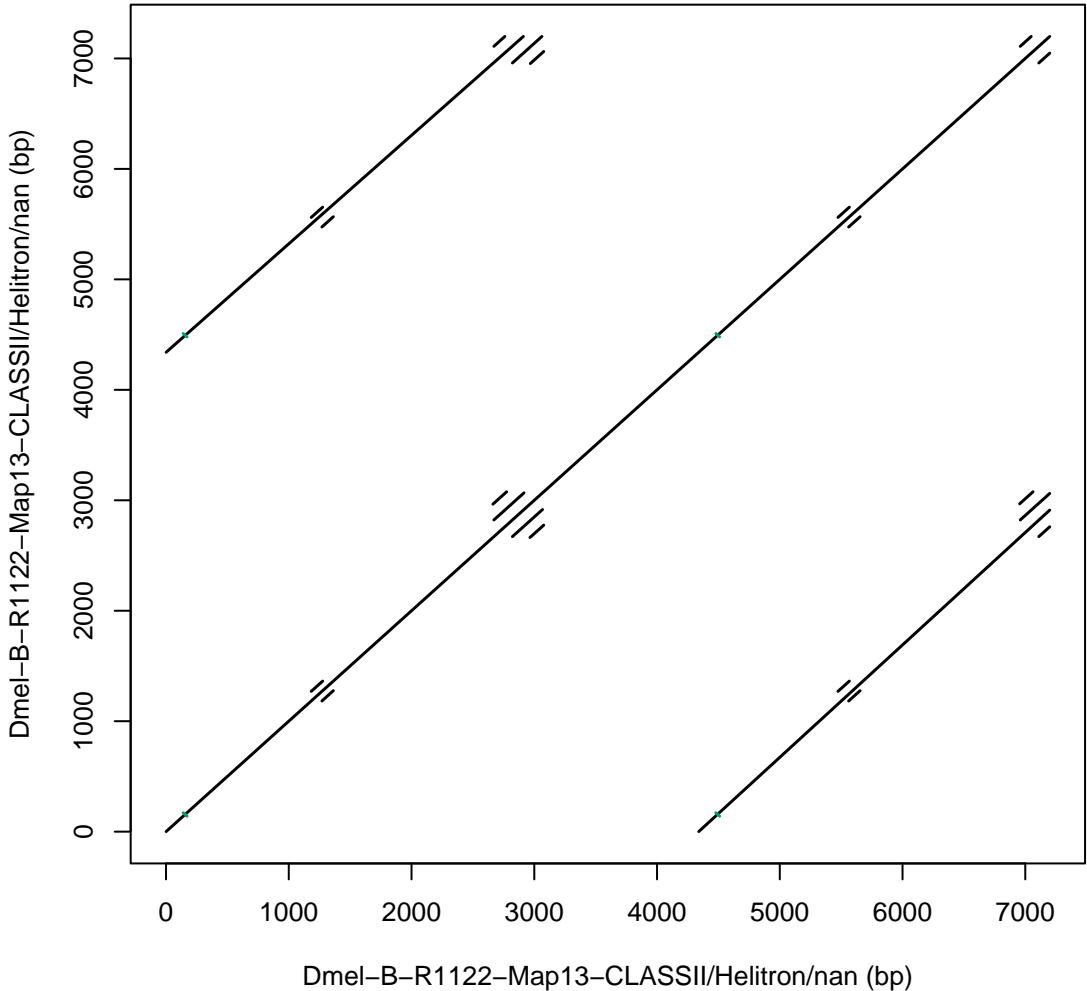
TE: Dmel-B-R1122-Map13-CLASSII/Helitron/nan
consensus size: 7199bp; fragments: 1481; full length: 3 (>=6479.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

