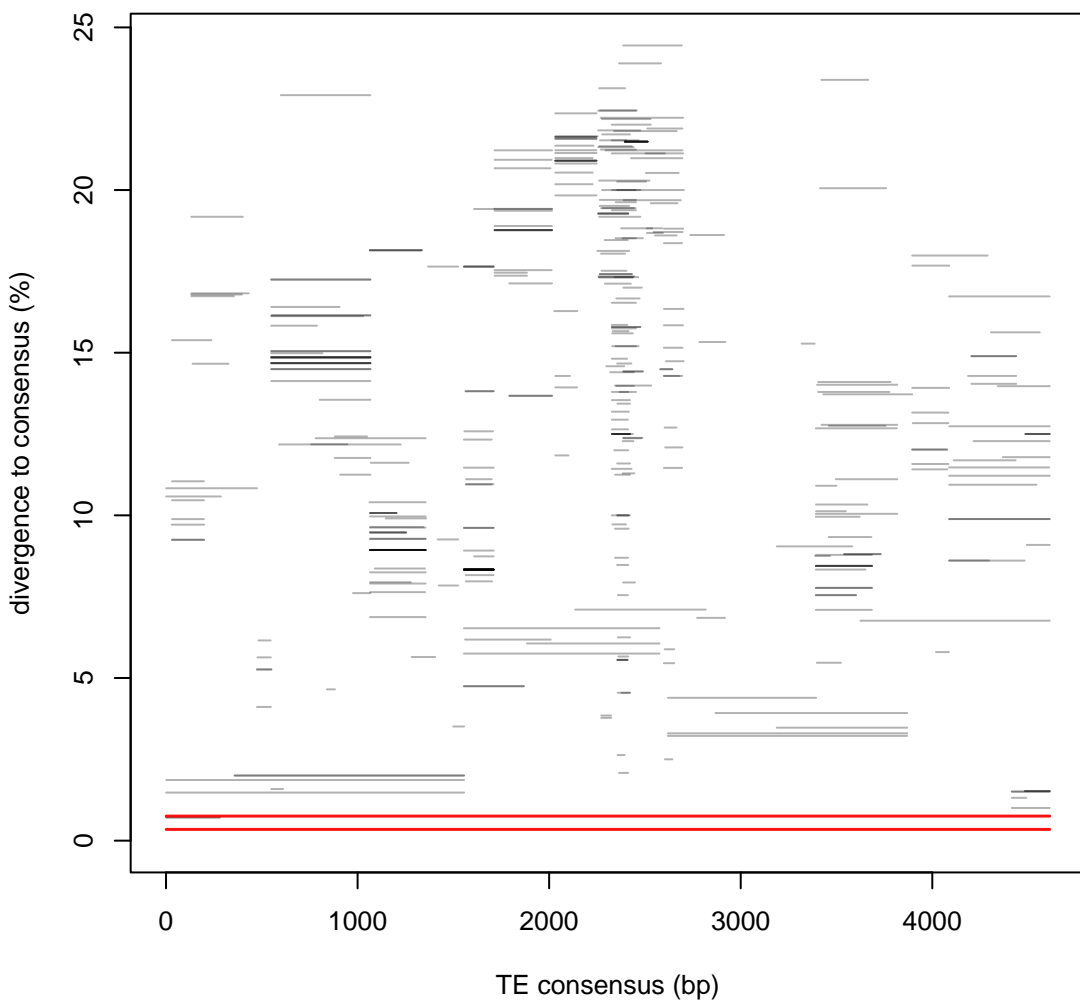
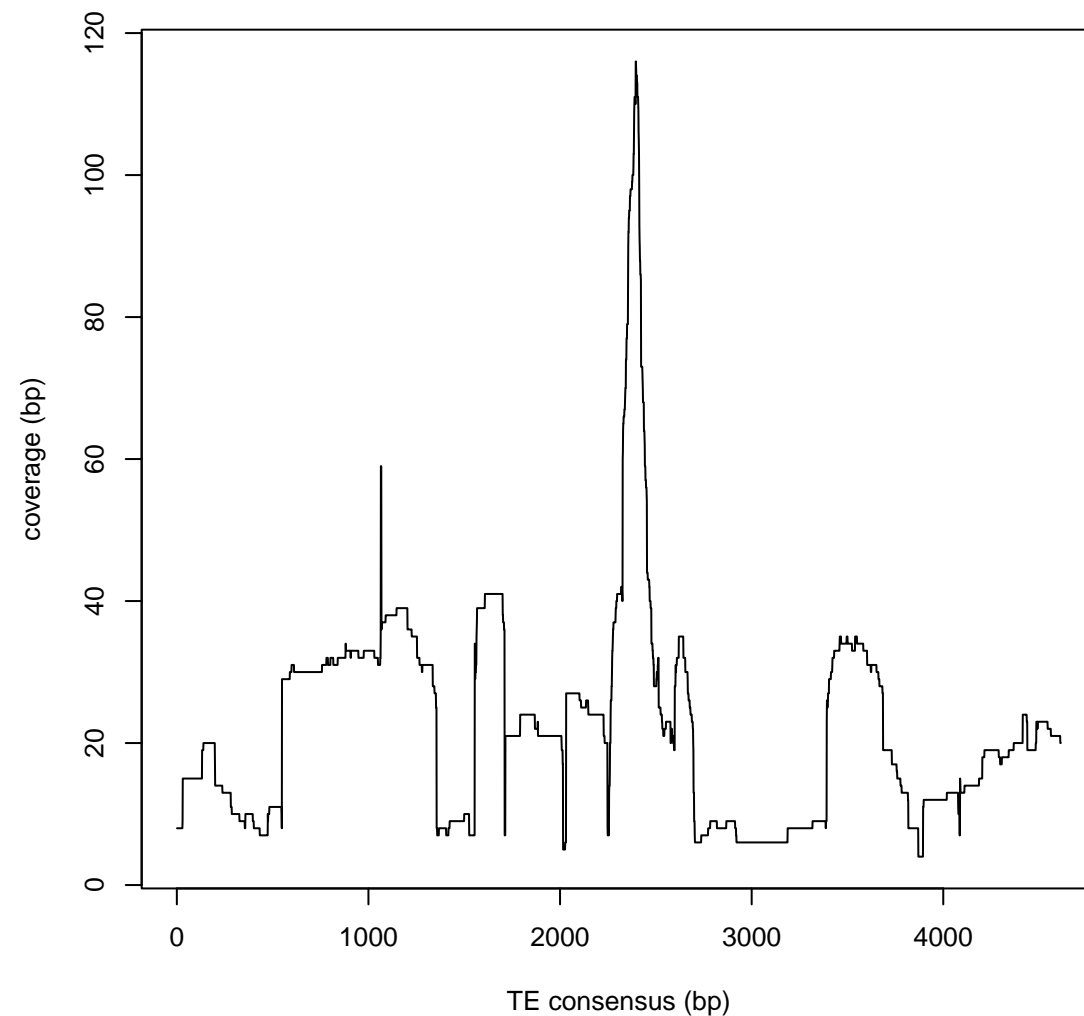


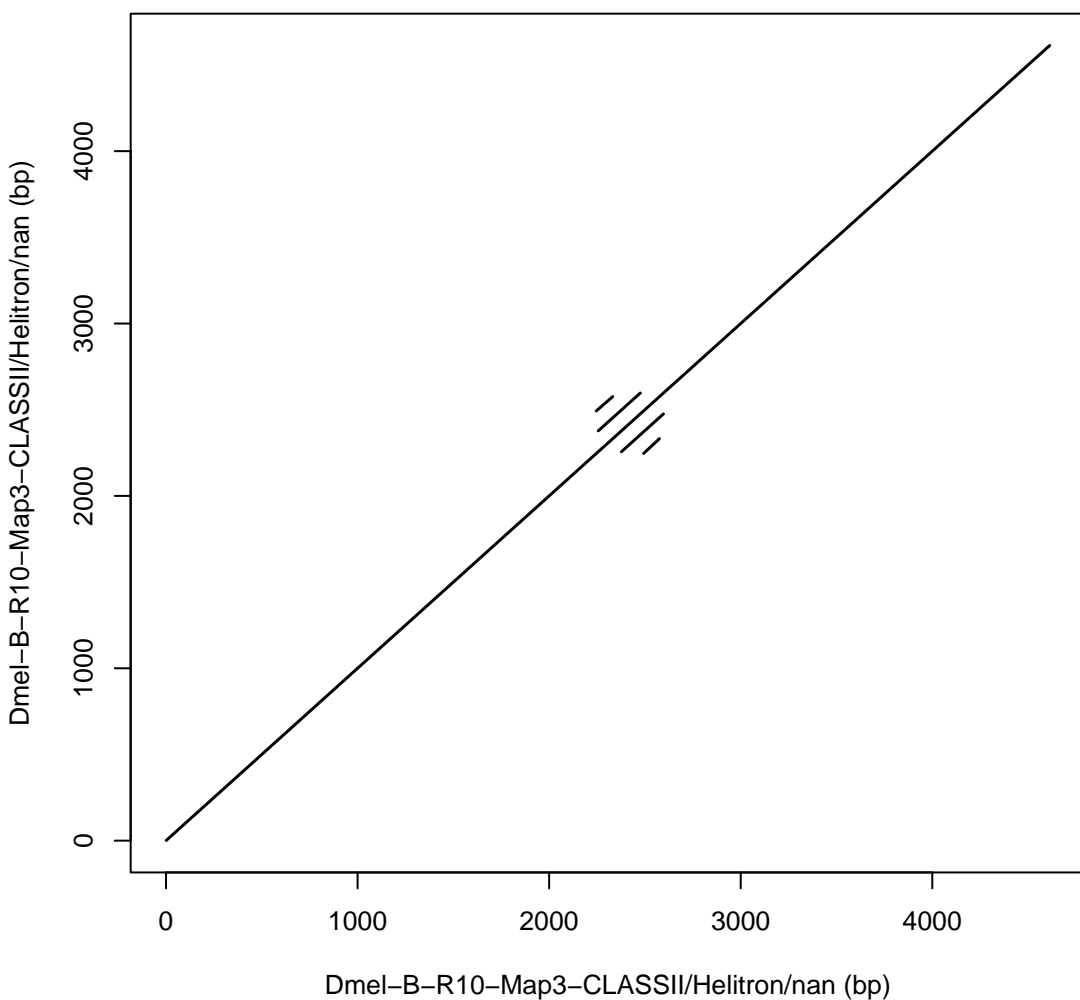
TE: Dmel-B-R10-Map3-CLASSII/Helitron/nan
consensus size: 4613bp; fragments: 395; full length: 2 (≥ 4151.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

