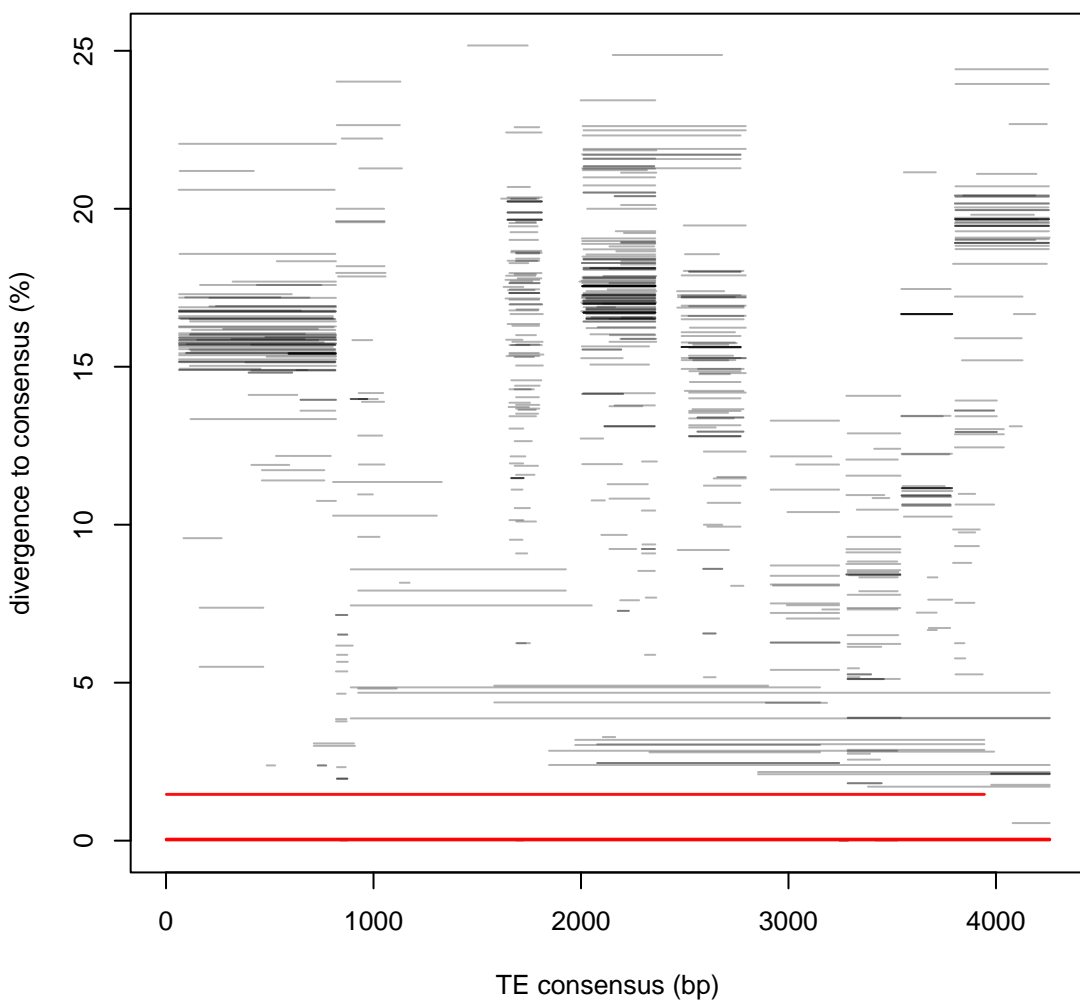
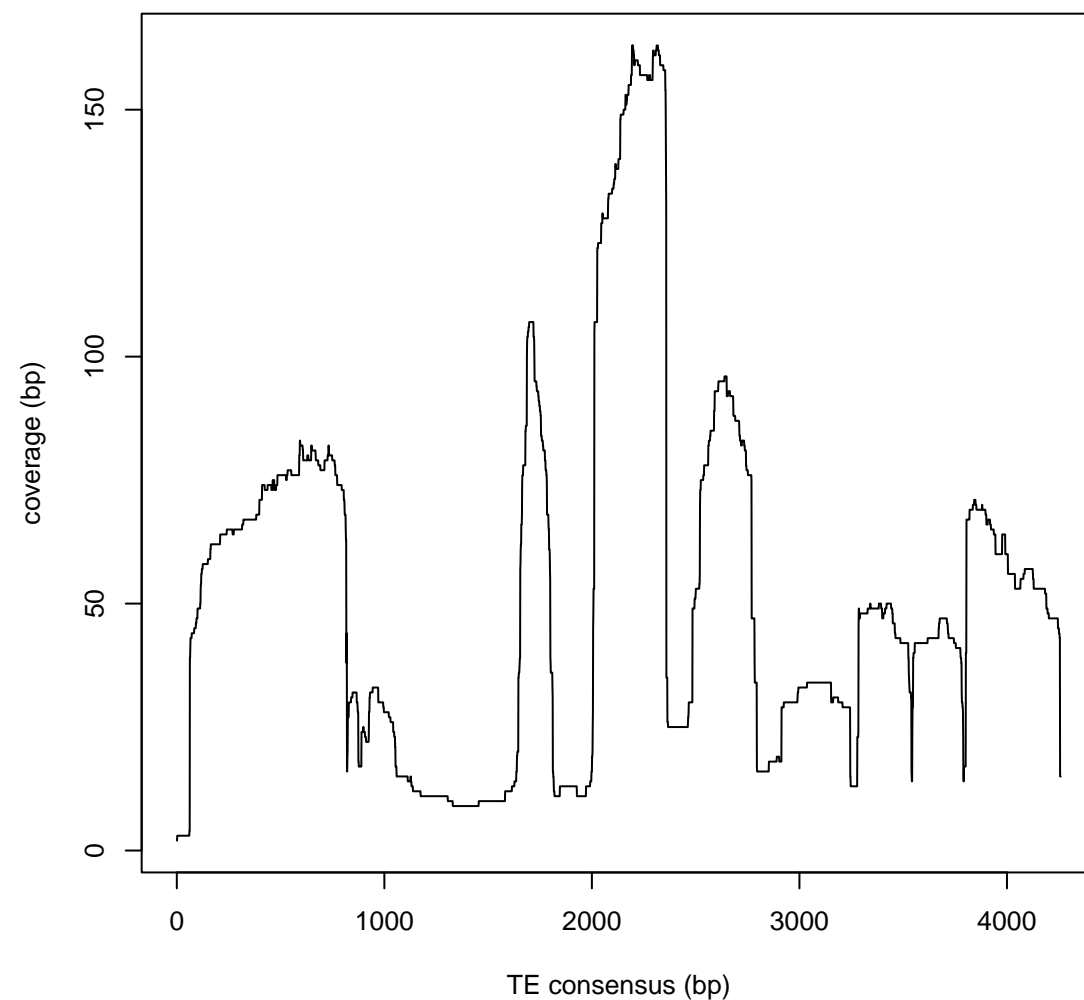


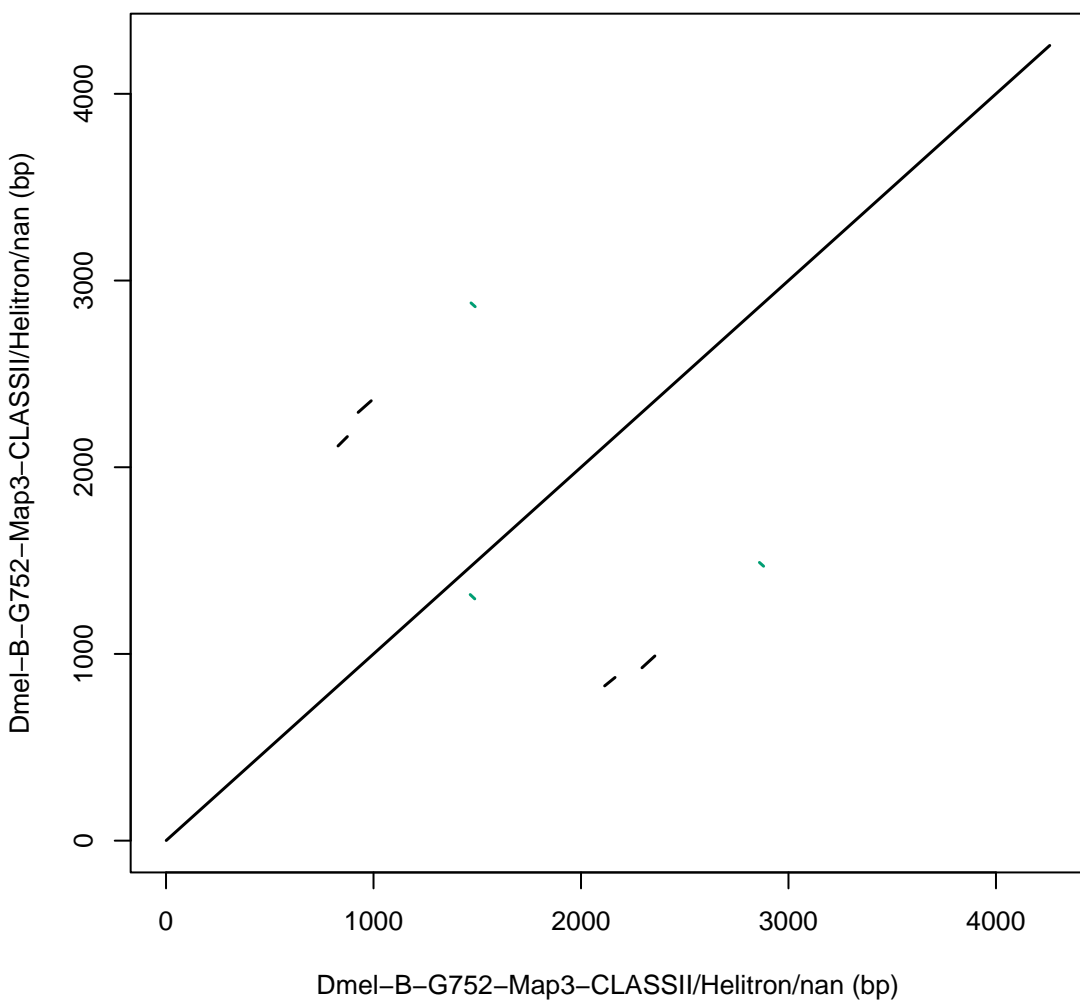
TE: Dmel-B-G752-Map3-CLASSII/Helitron/nan
consensus size: 4259bp; fragments: 655; full length: 3 (≥ 3833.1 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

