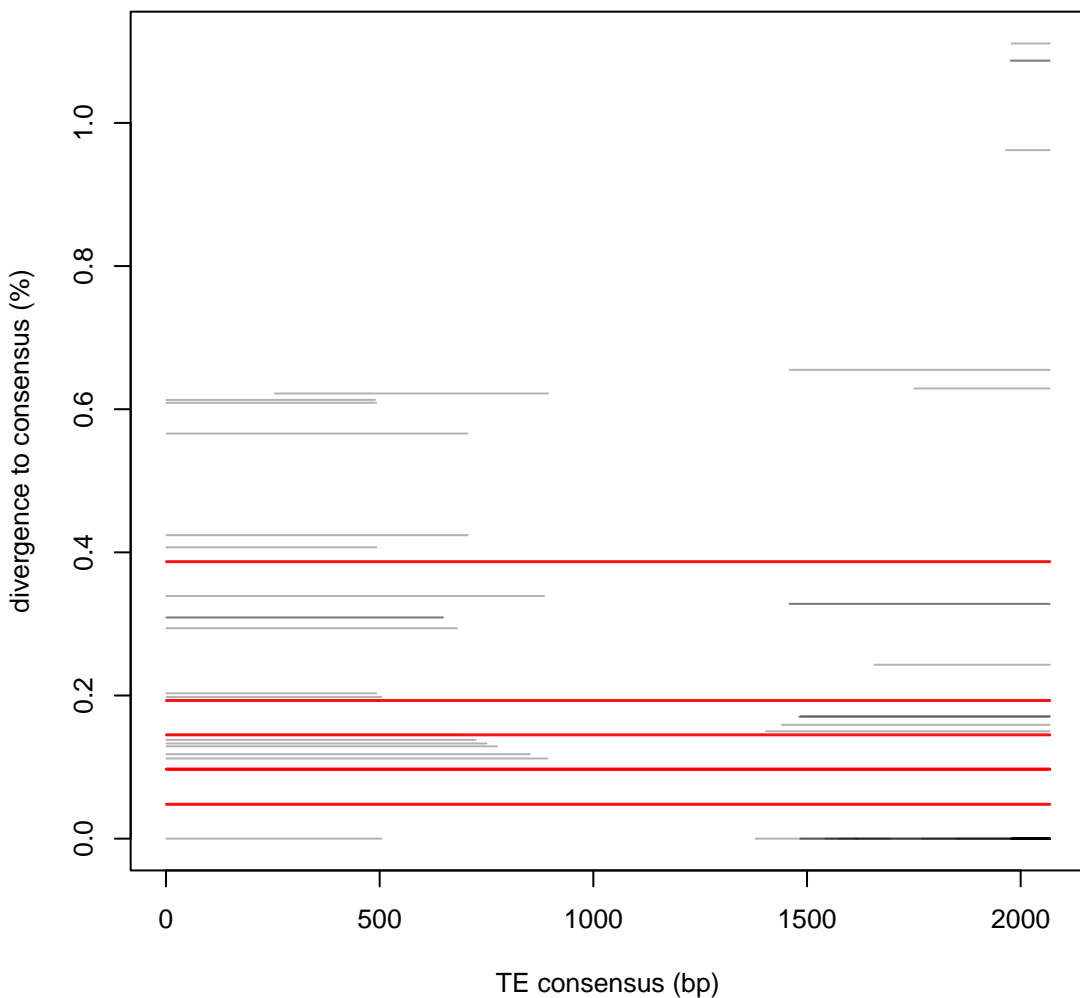
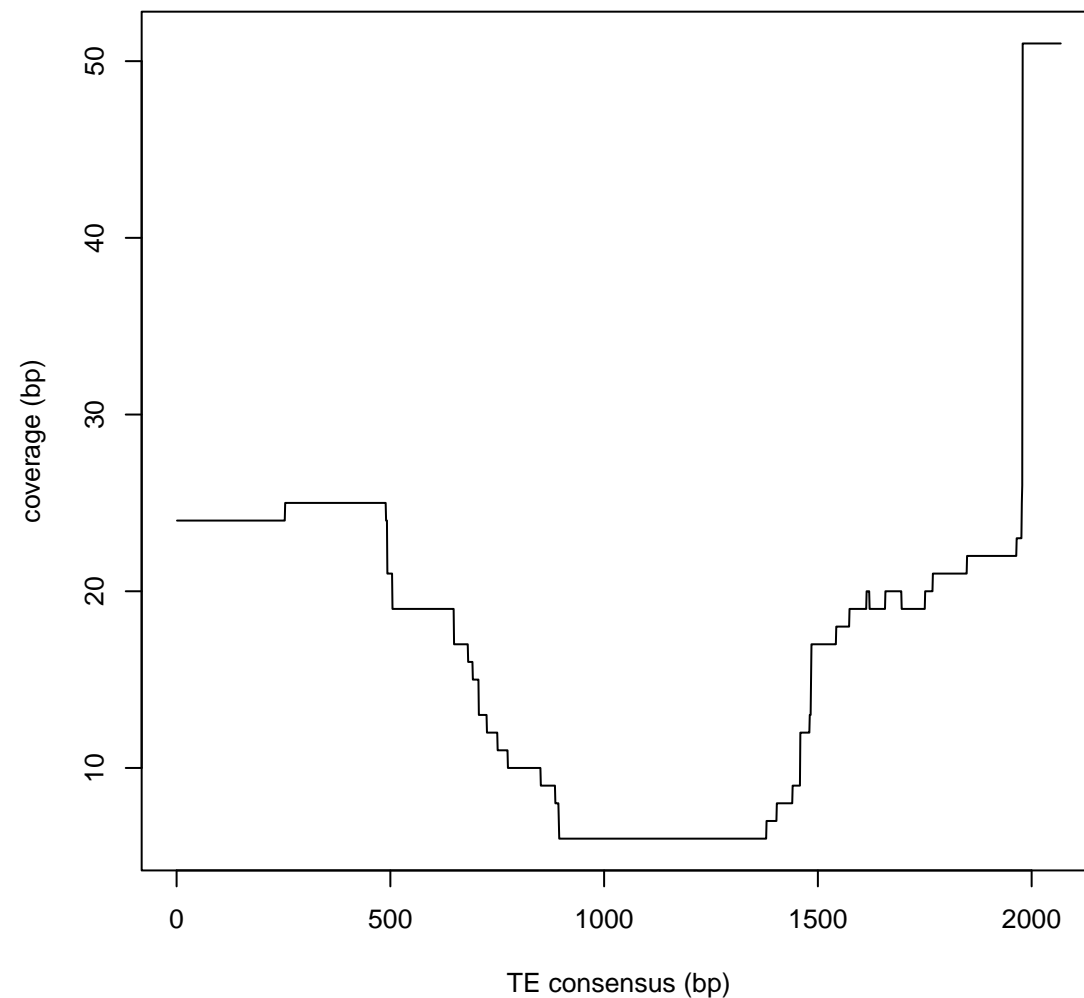


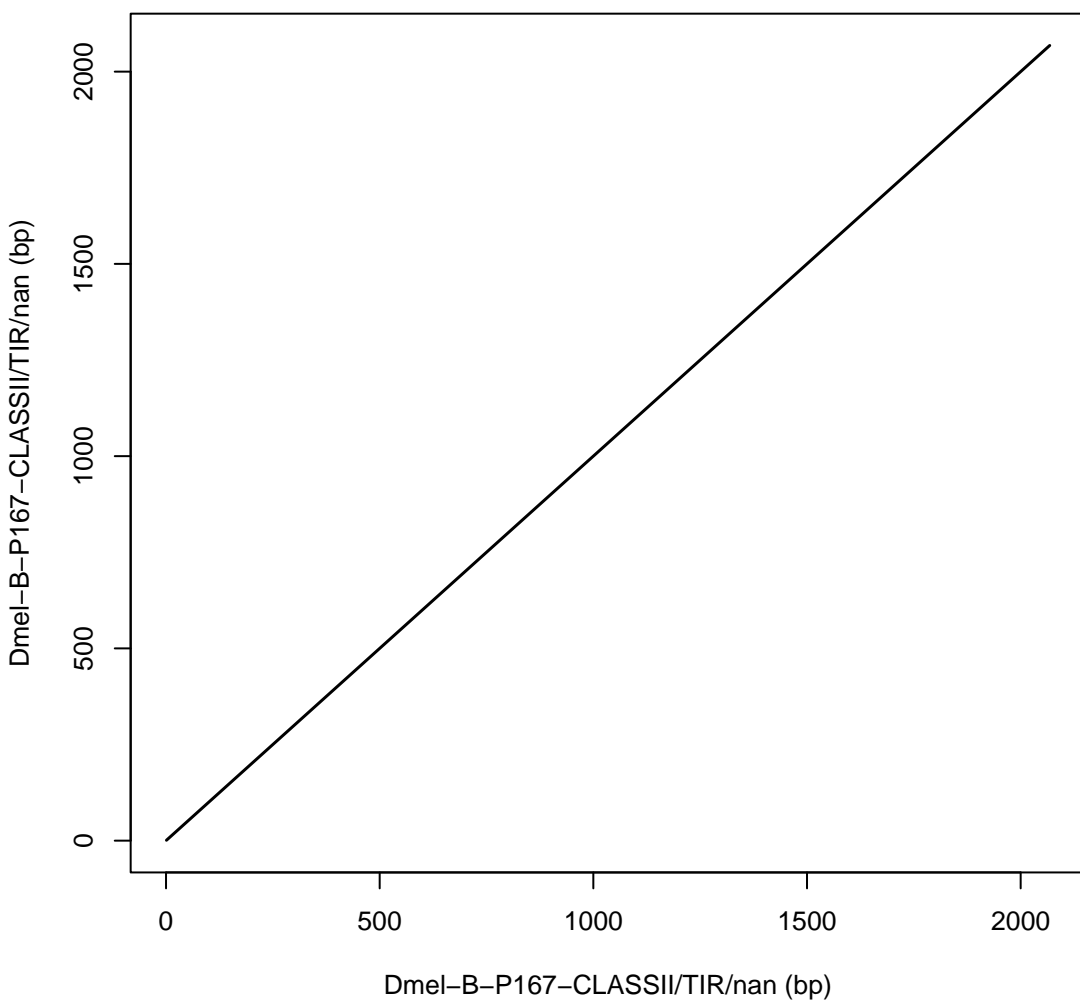
TE: Dmel-B-P167-CLASSII/TIR/nan
consensus size: 2068bp; fragments: 72; full length: 6 ($\geq 1861.2\text{bp}$)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

