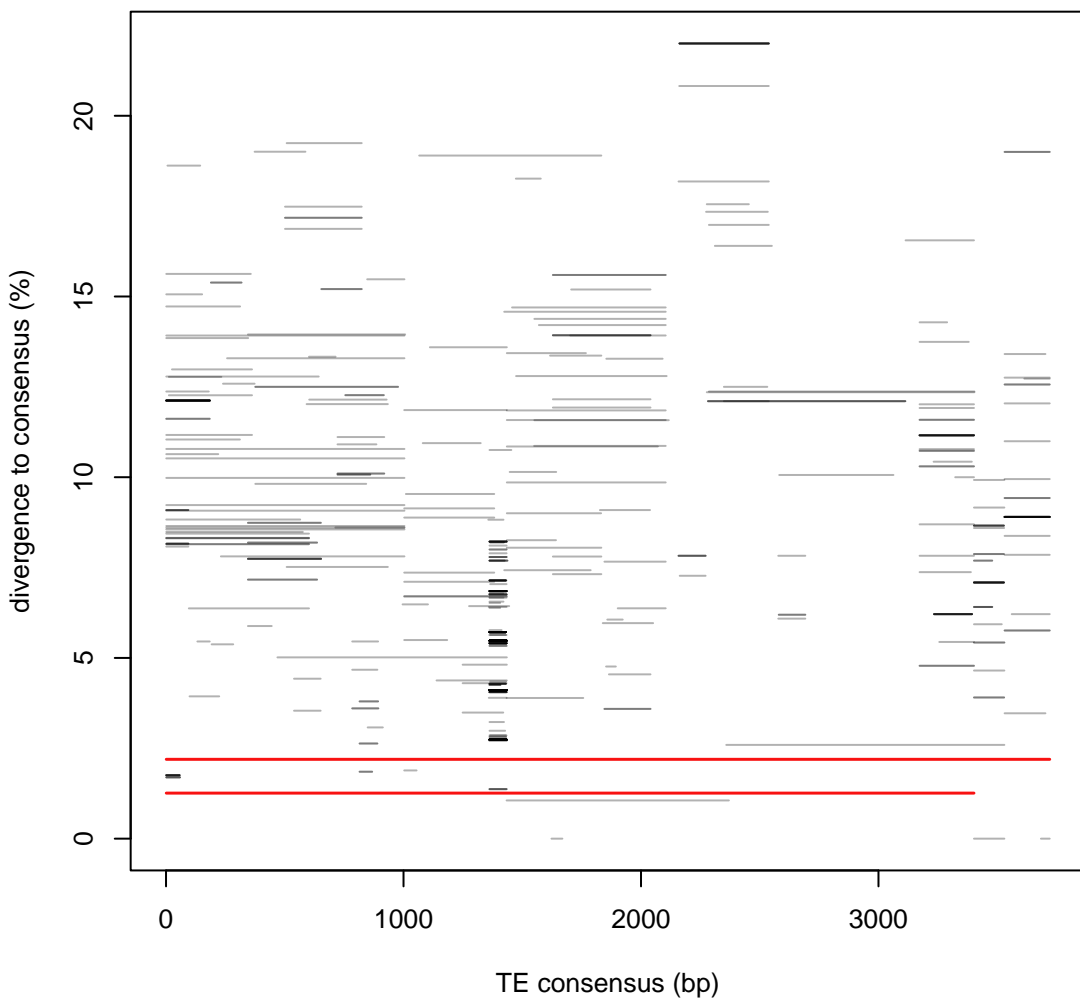
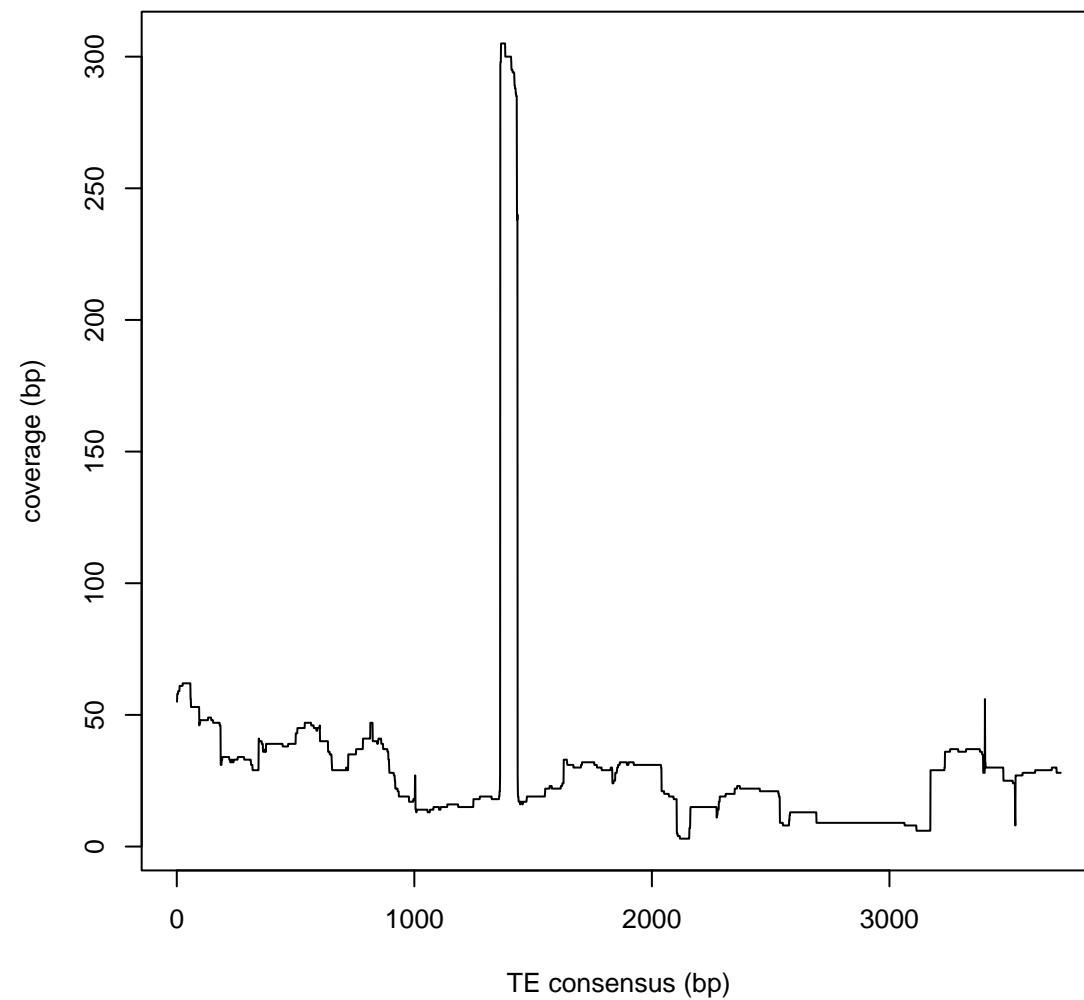


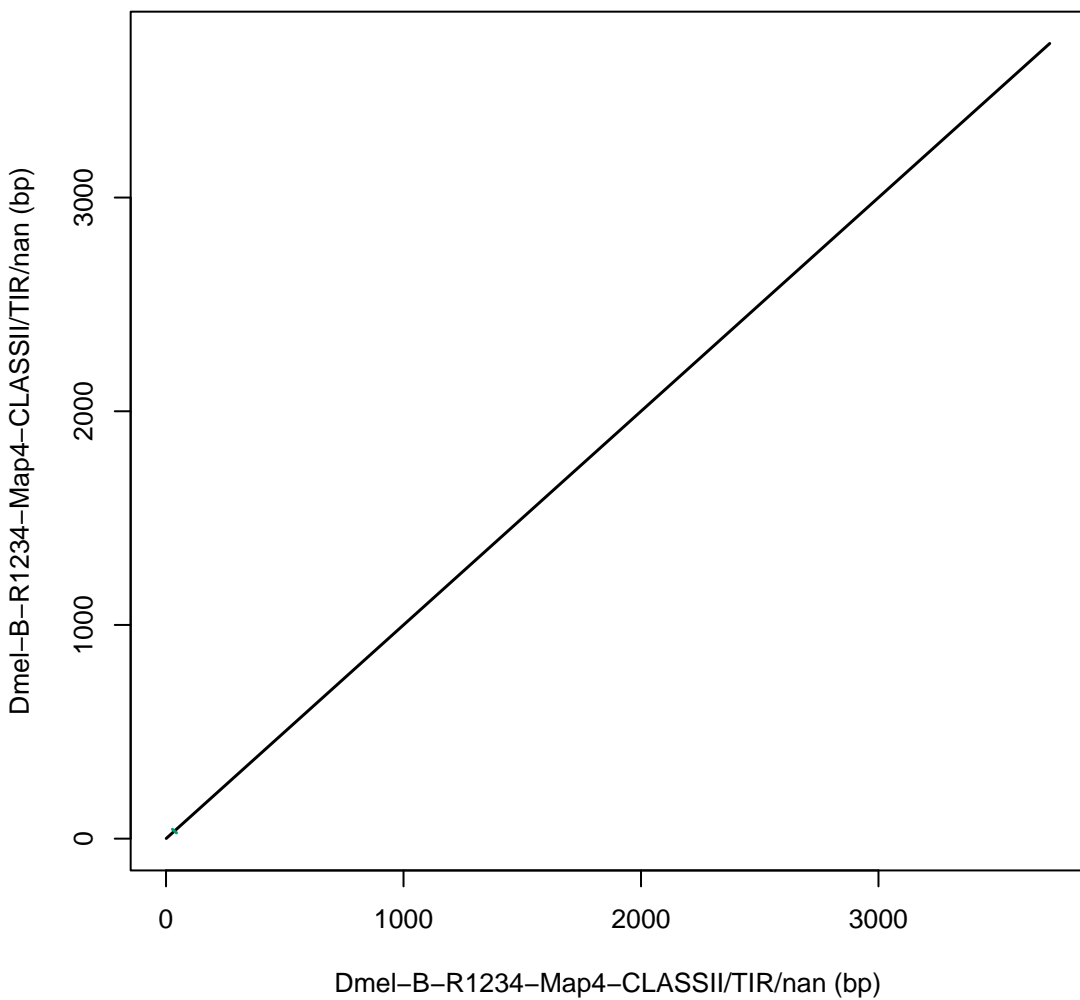
TE: Dmel-B-R1234-Map4-CLASSII/TIR/nan  
consensus size: 3721bp; fragments: 590; full length: 2 (>=3348.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

