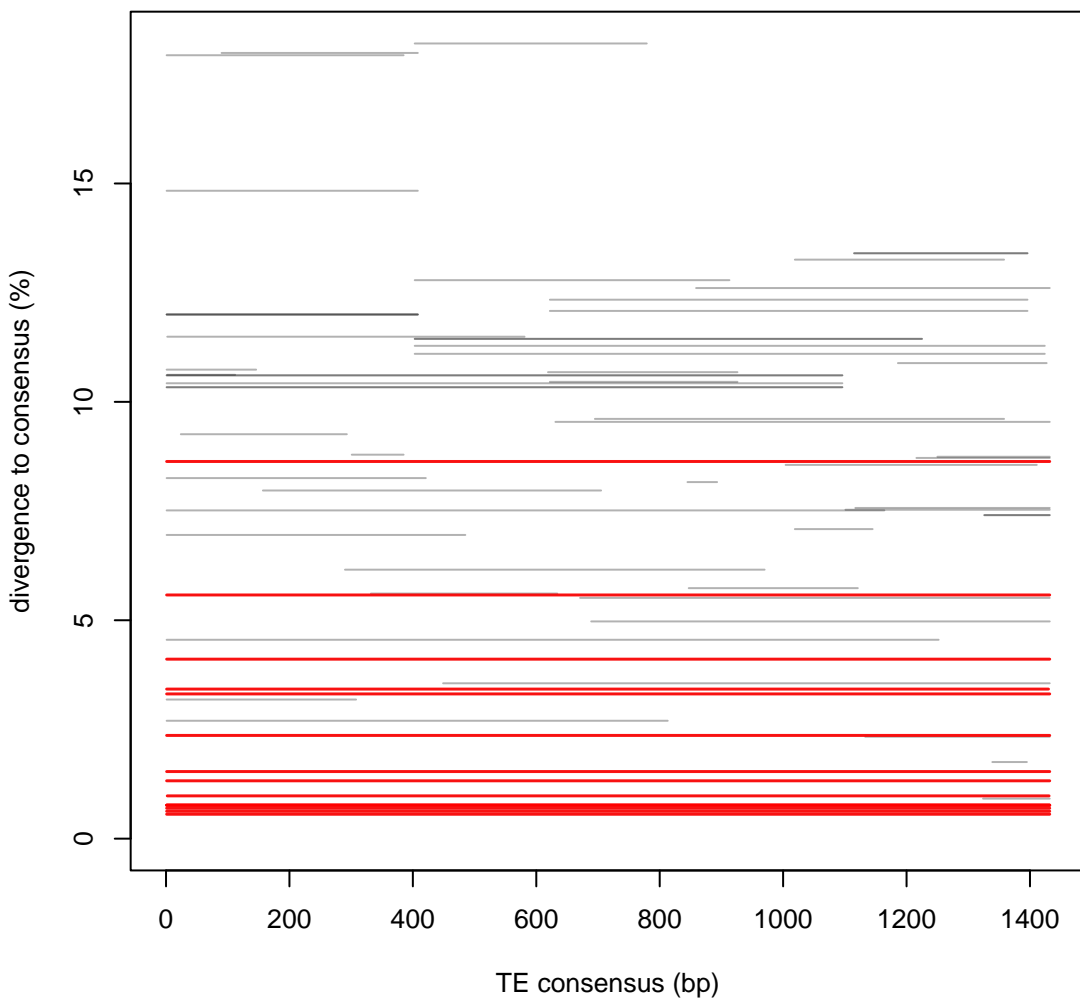
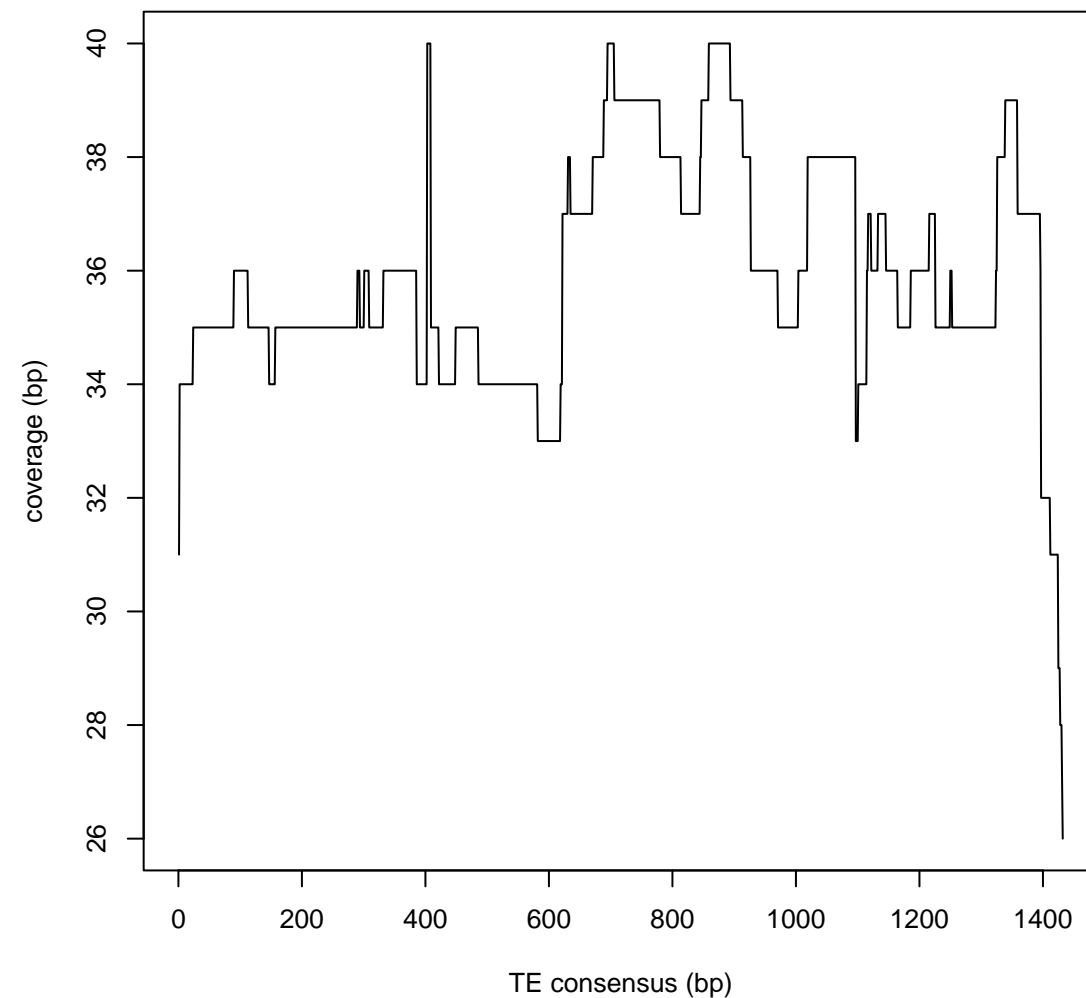


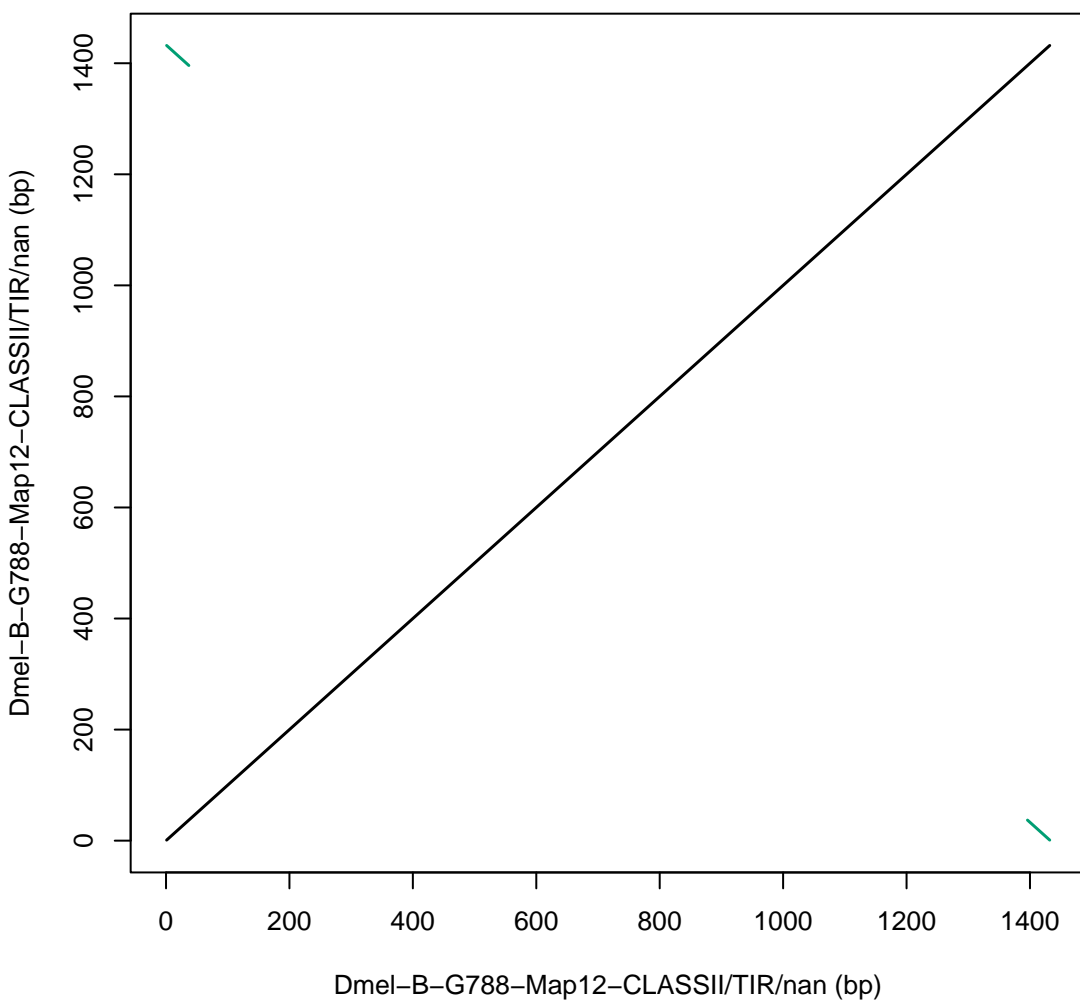
TE: Dmel-B-G788-Map12-CLASSII/TIR/nan  
consensus size: 1432bp; fragments: 73; full length: 15 ( $\geq 1288.8$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

