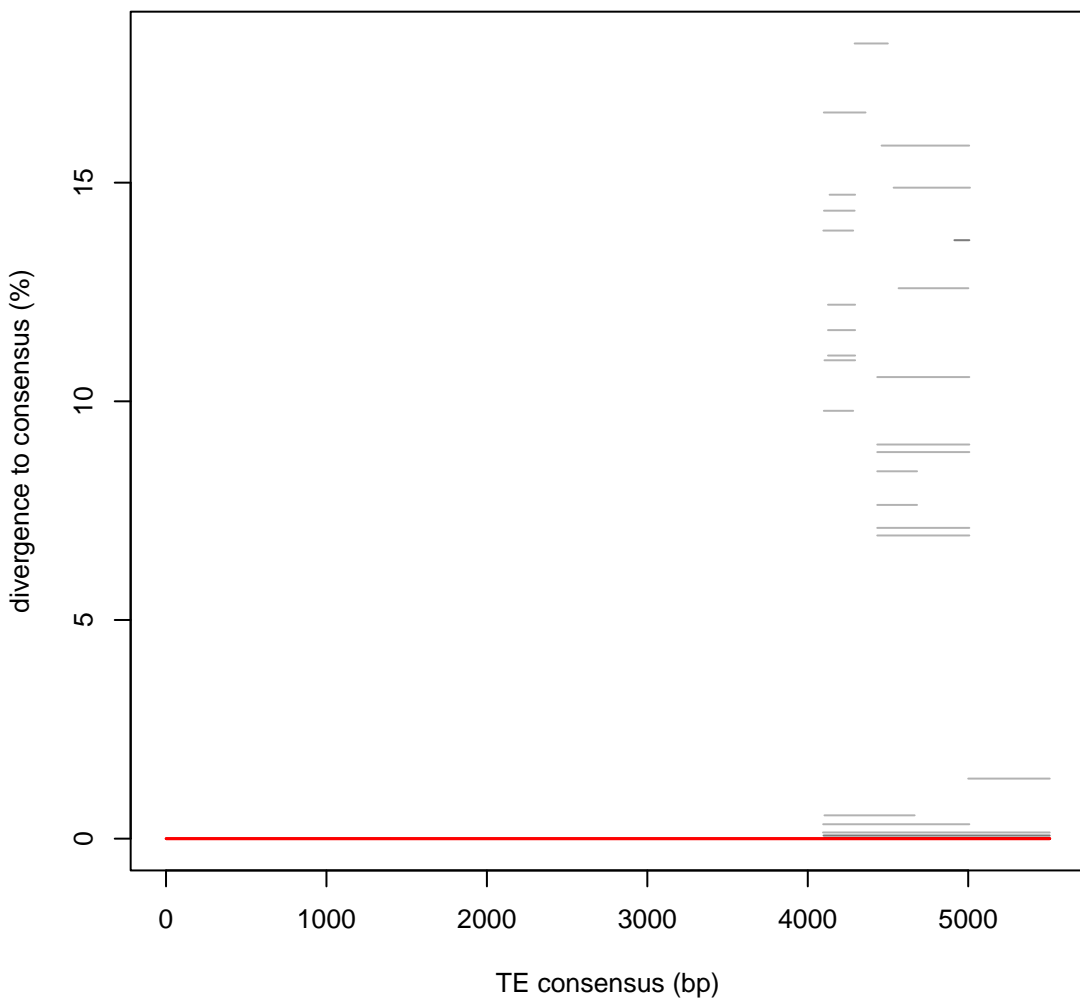
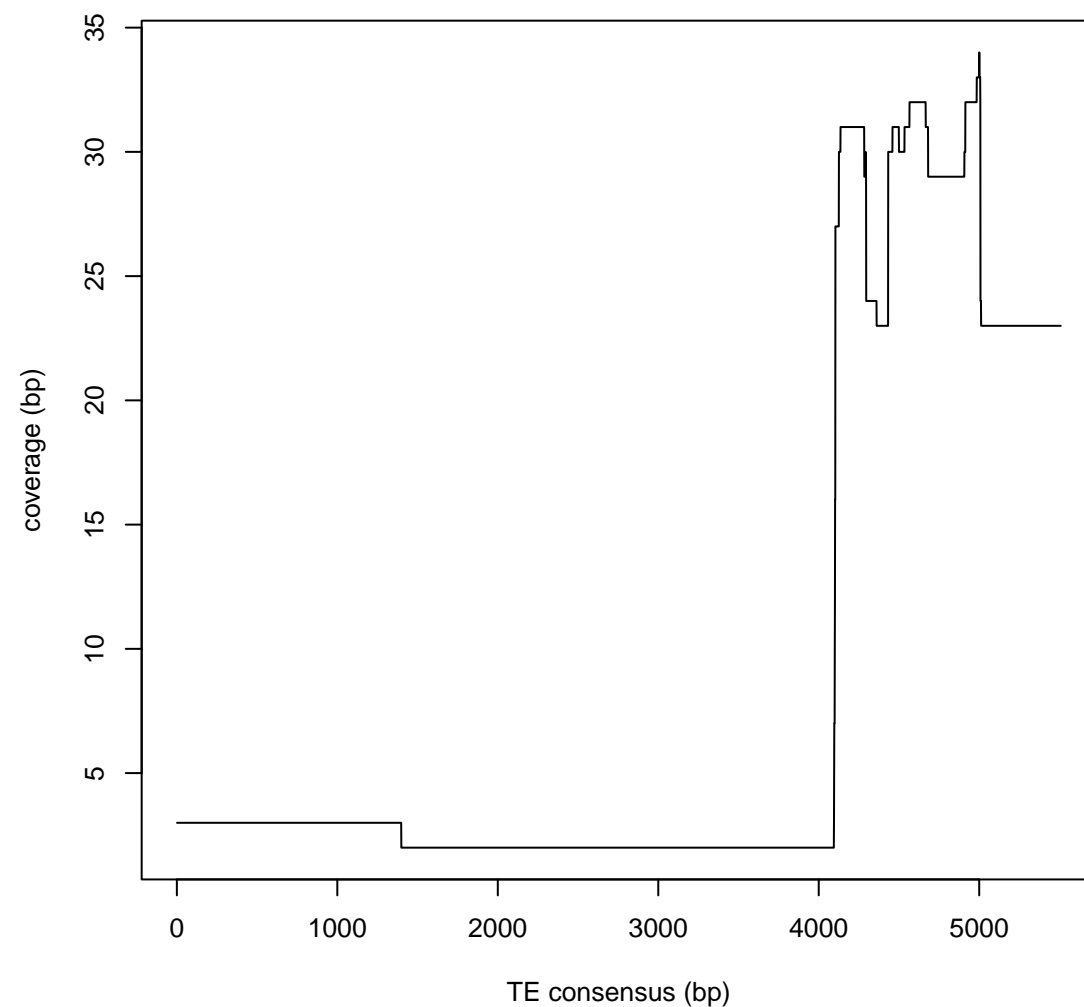


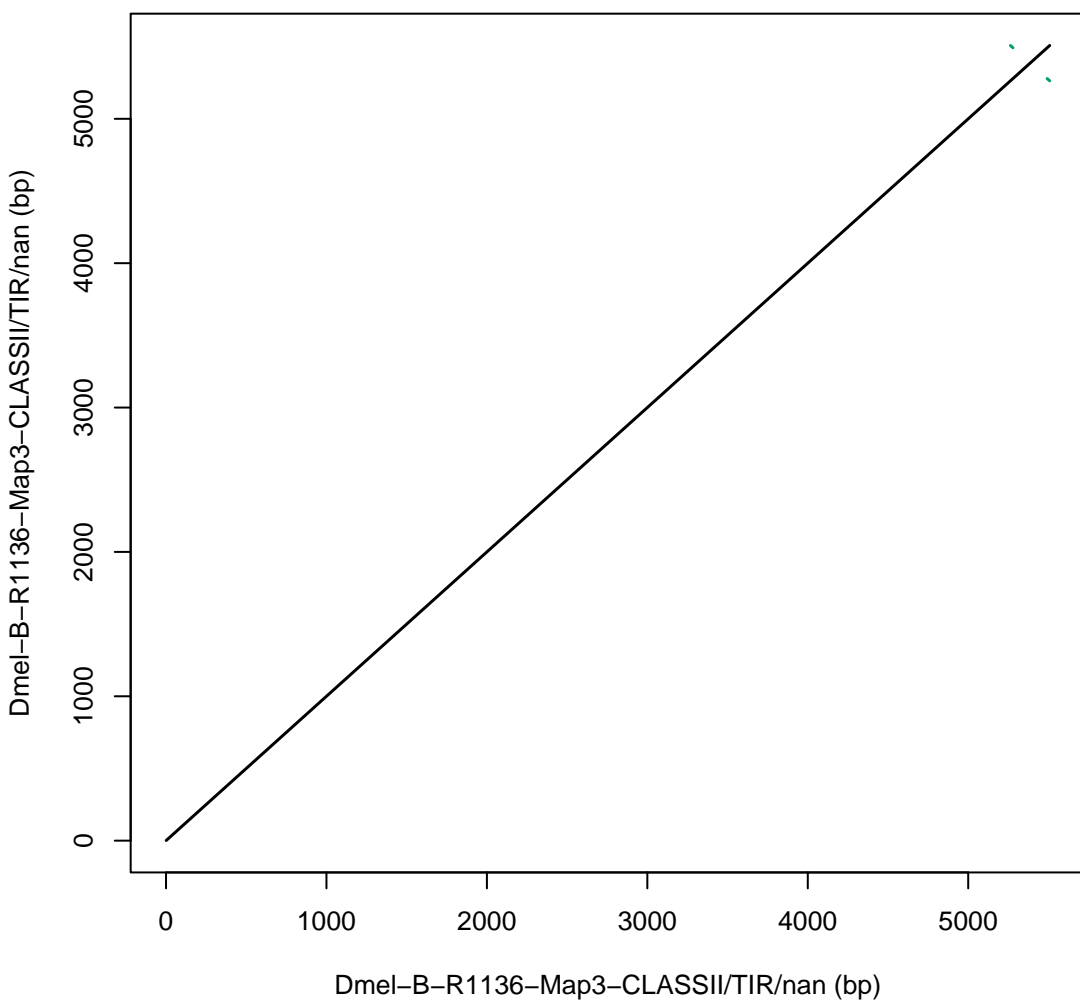
TE: Dmel-B-R1136-Map3-CLASSII/TIR/nan  
consensus size: 5508bp; fragments: 48; full length: 2 (>=4957.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

