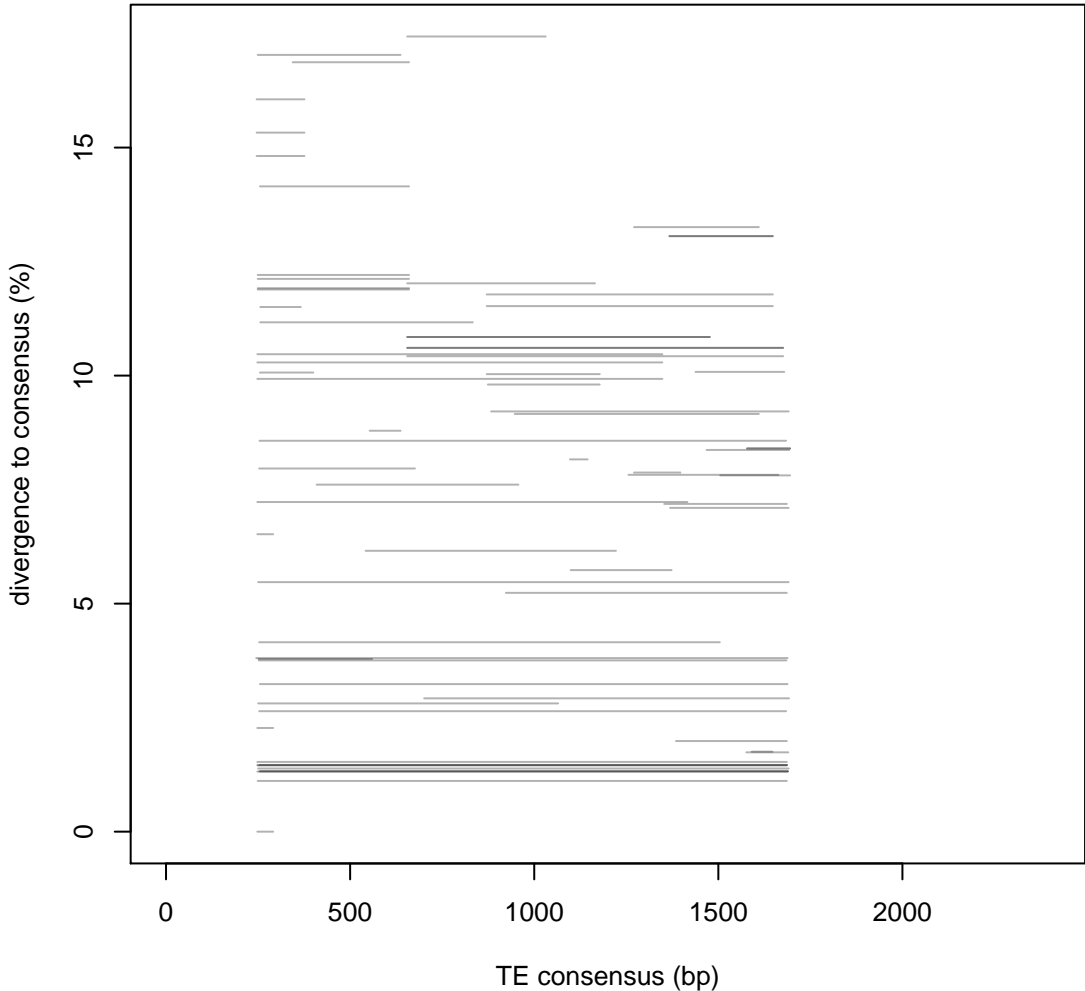
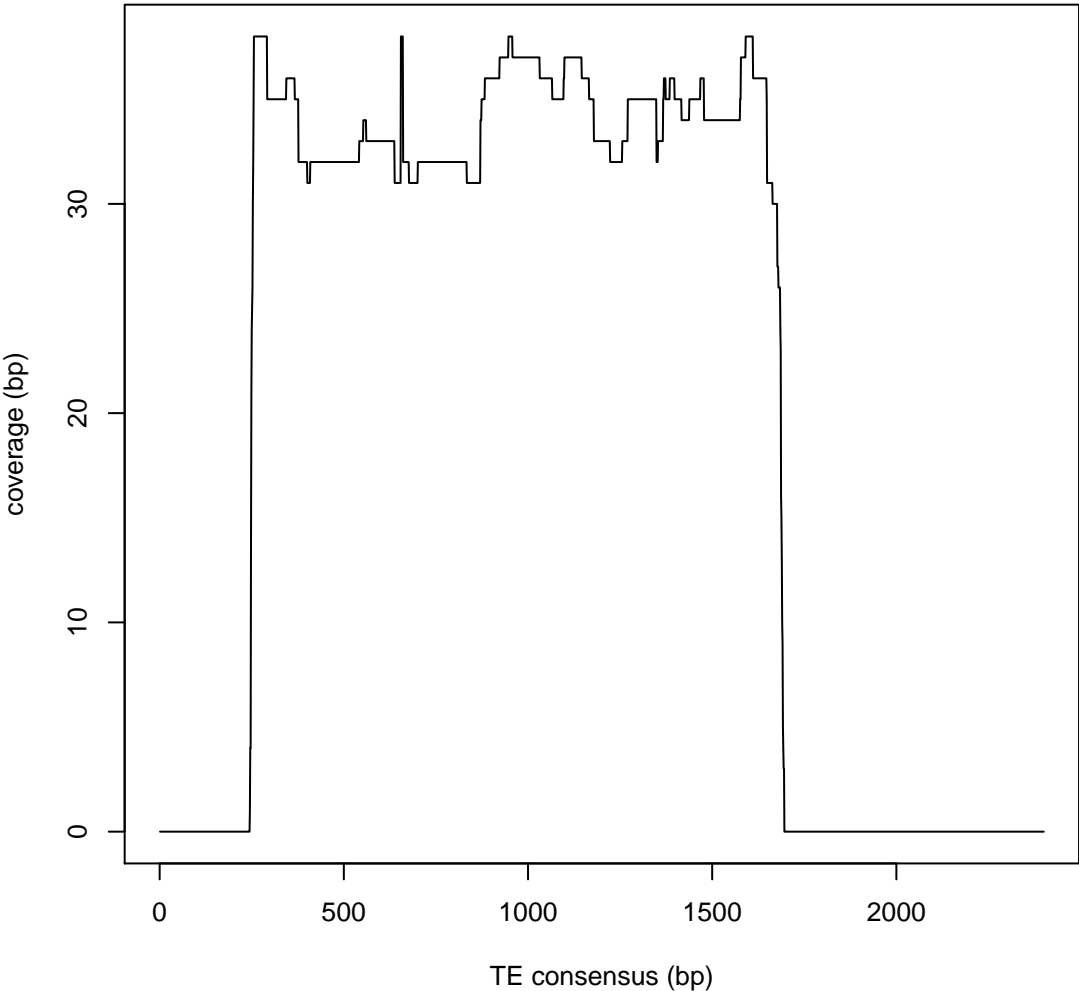


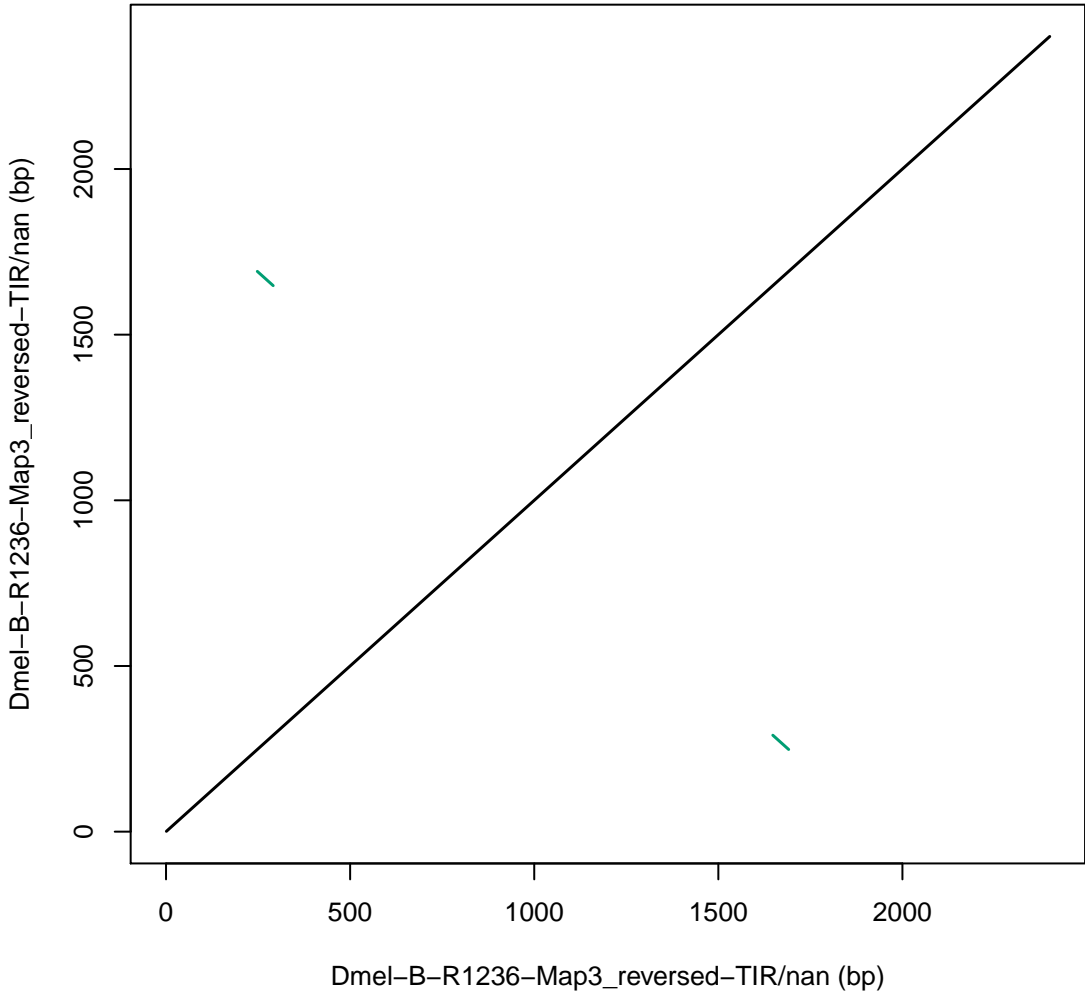
TE: Dmel-B-R1236-Map3\_reversed-TIR/nan  
consensus size: 2400bp; fragments: 74; full length: 0 (>=2160bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

