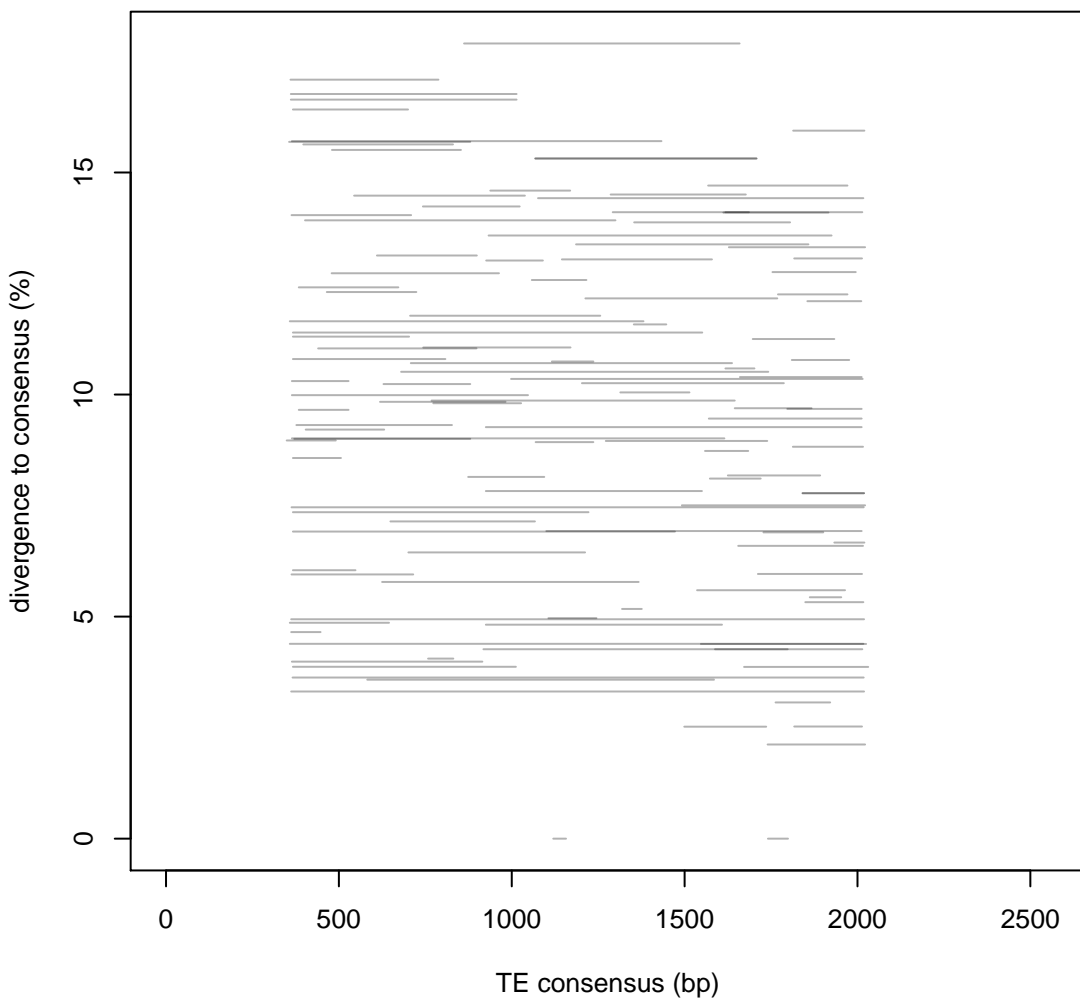
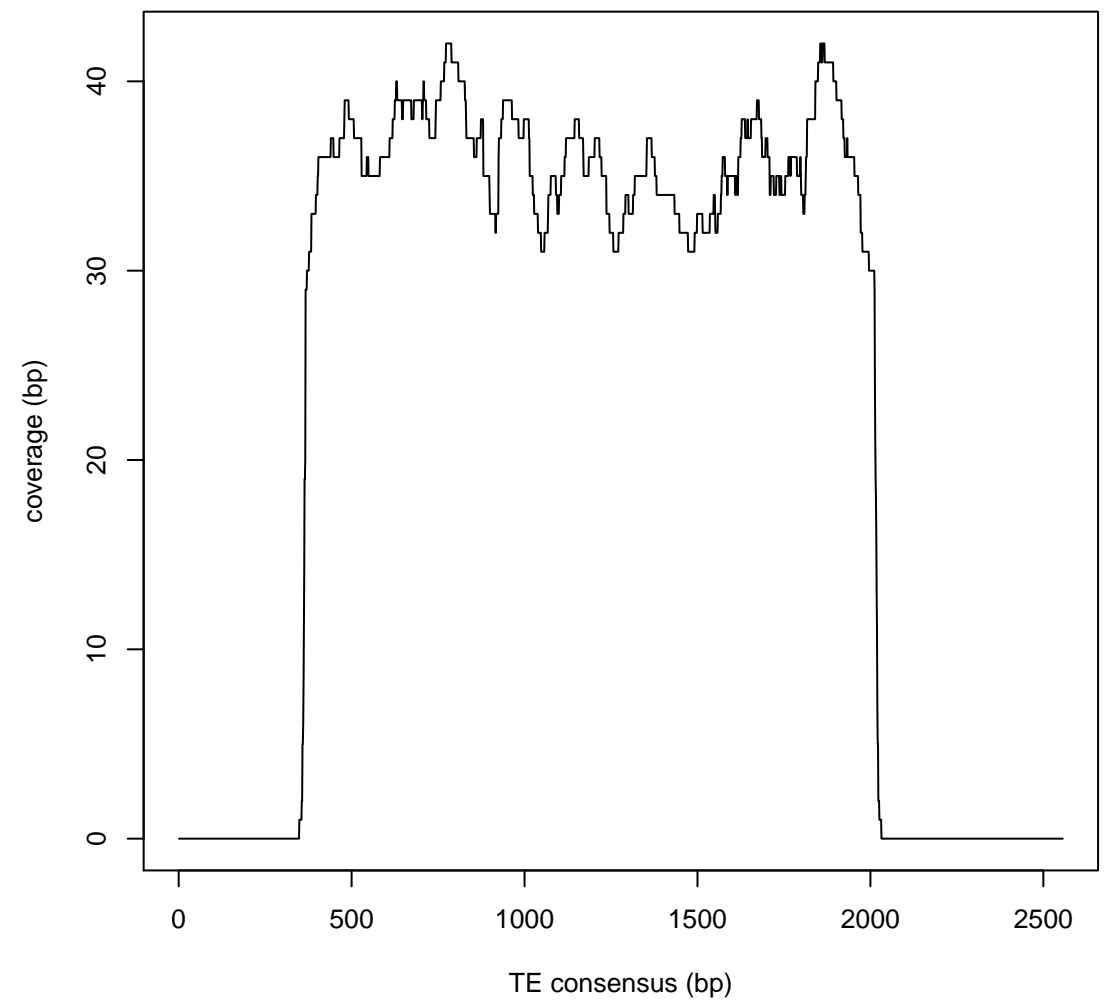


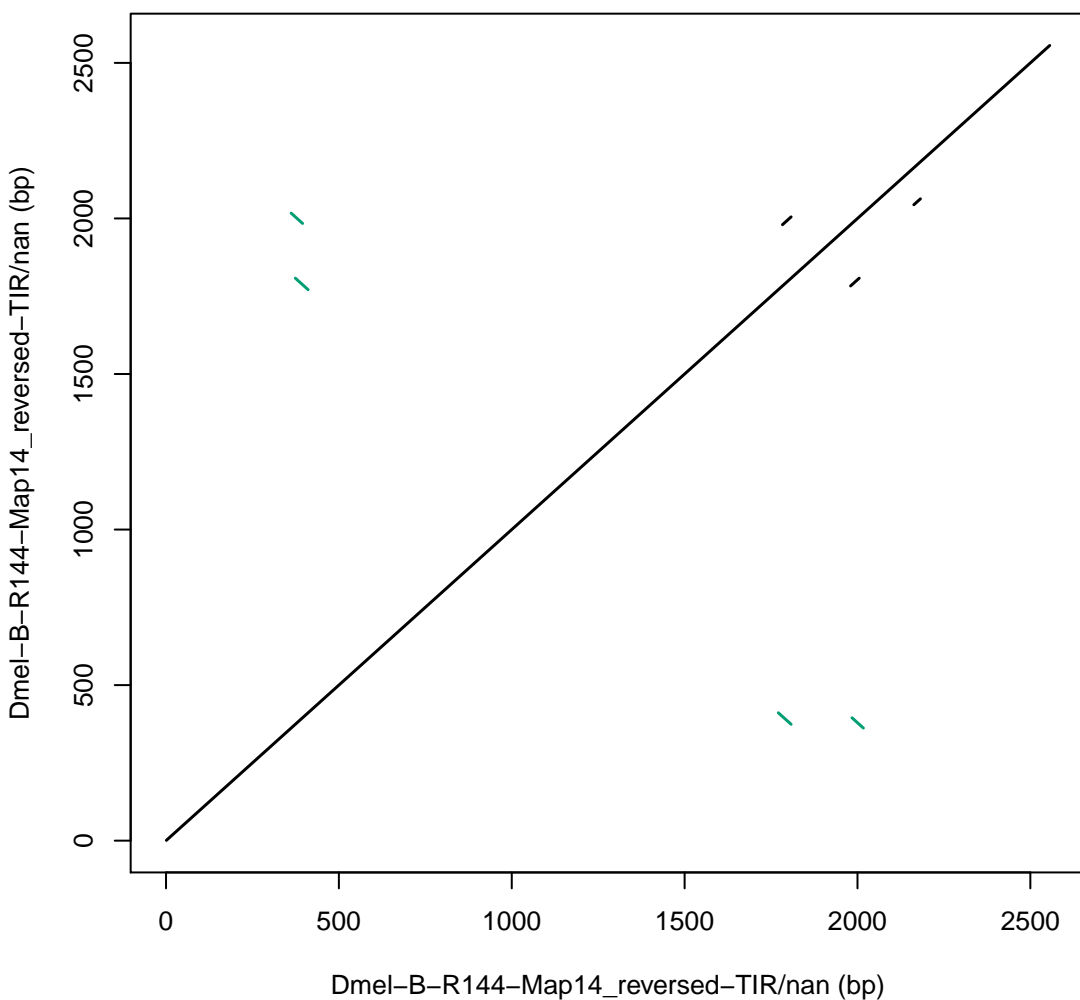
TE: Dmel-B-R144-Map14\_reversed-TIR/nan  
consensus size: 2556bp; fragments: 124; full length: 0 (>=2300.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

