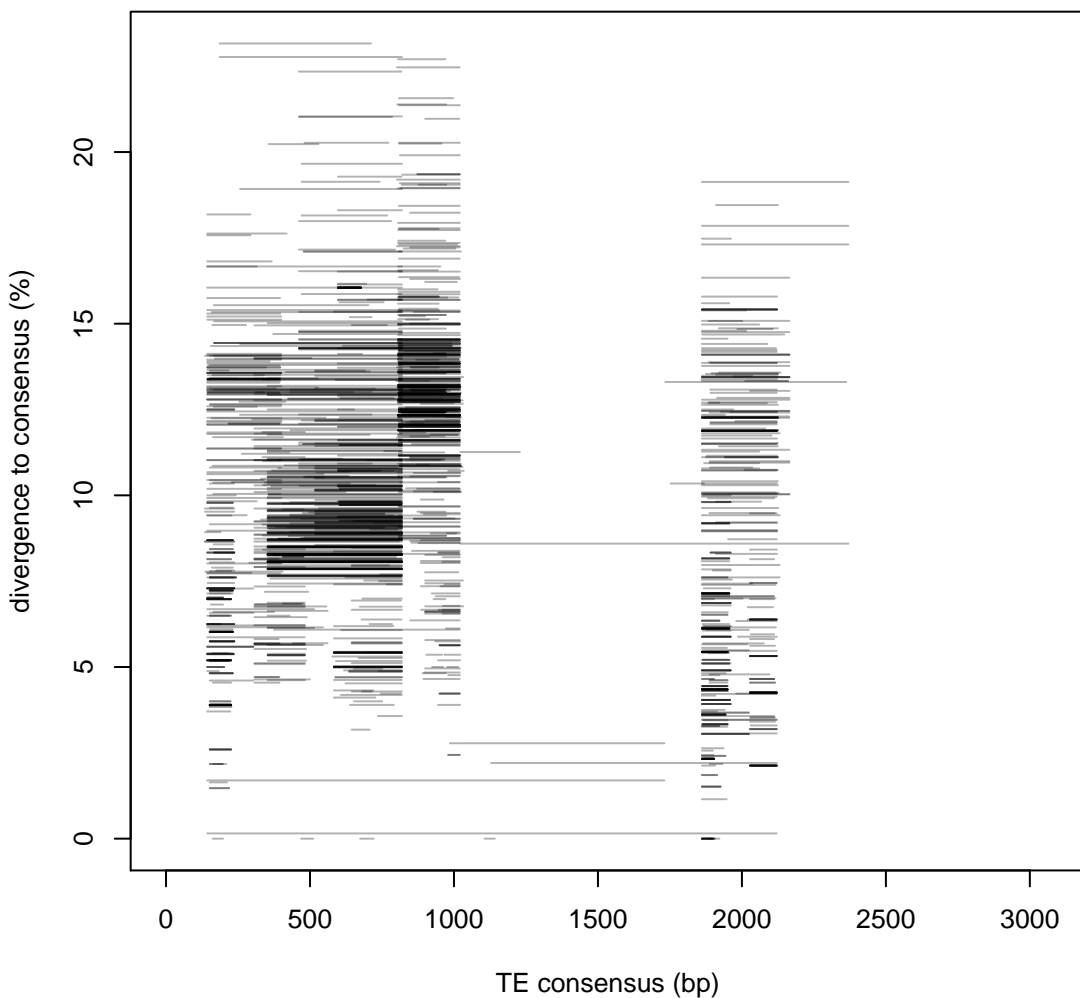
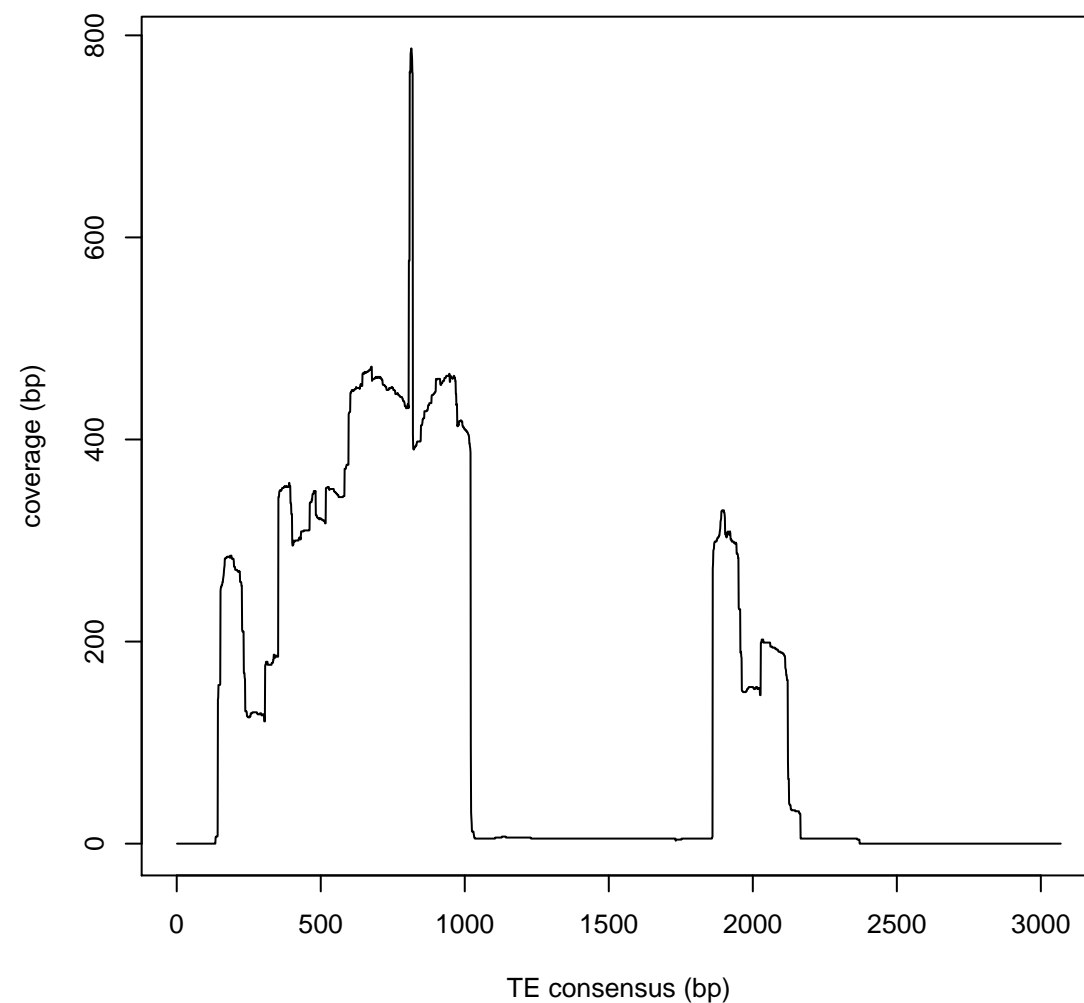


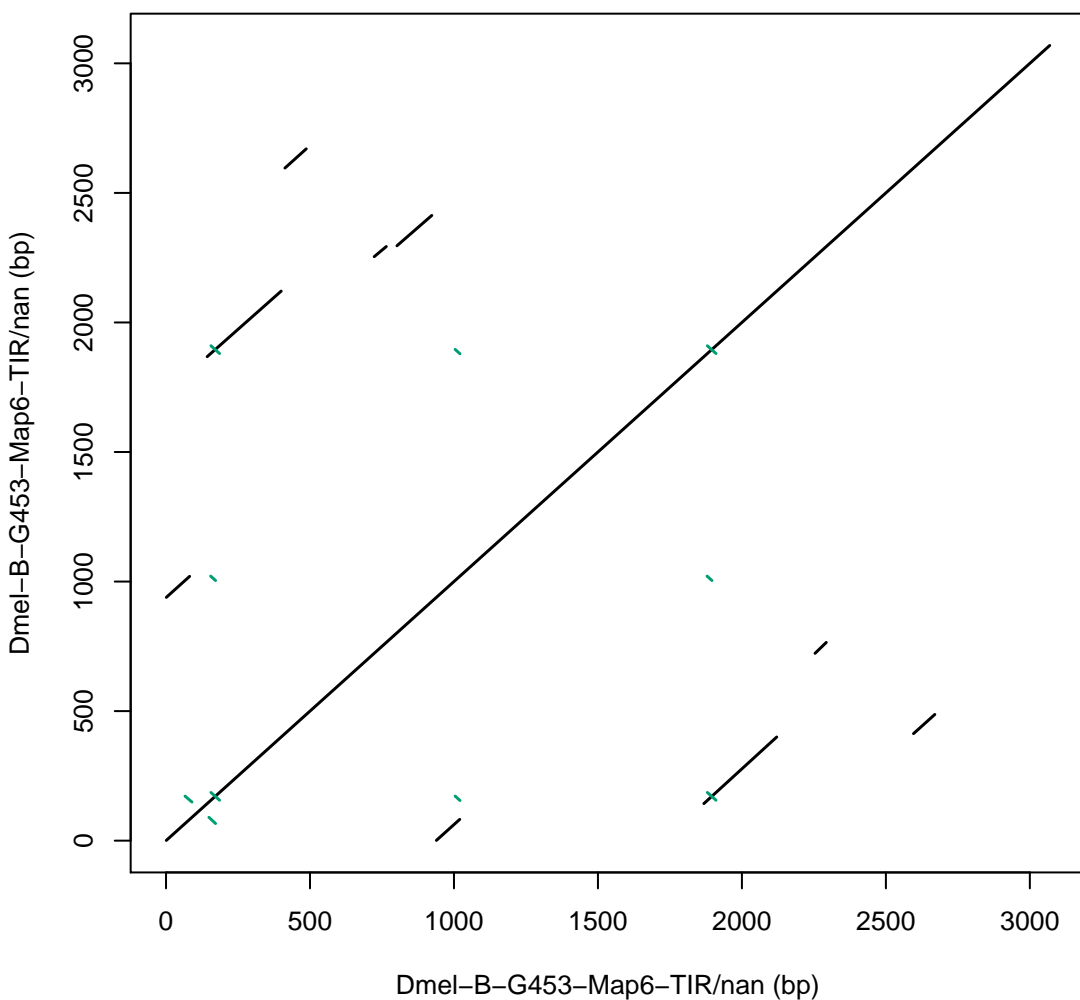
TE: Dmel-B-G453-Map6-TIR/nan  
consensus size: 3069bp; fragments: 1712; full length: 0 (>=2762.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

