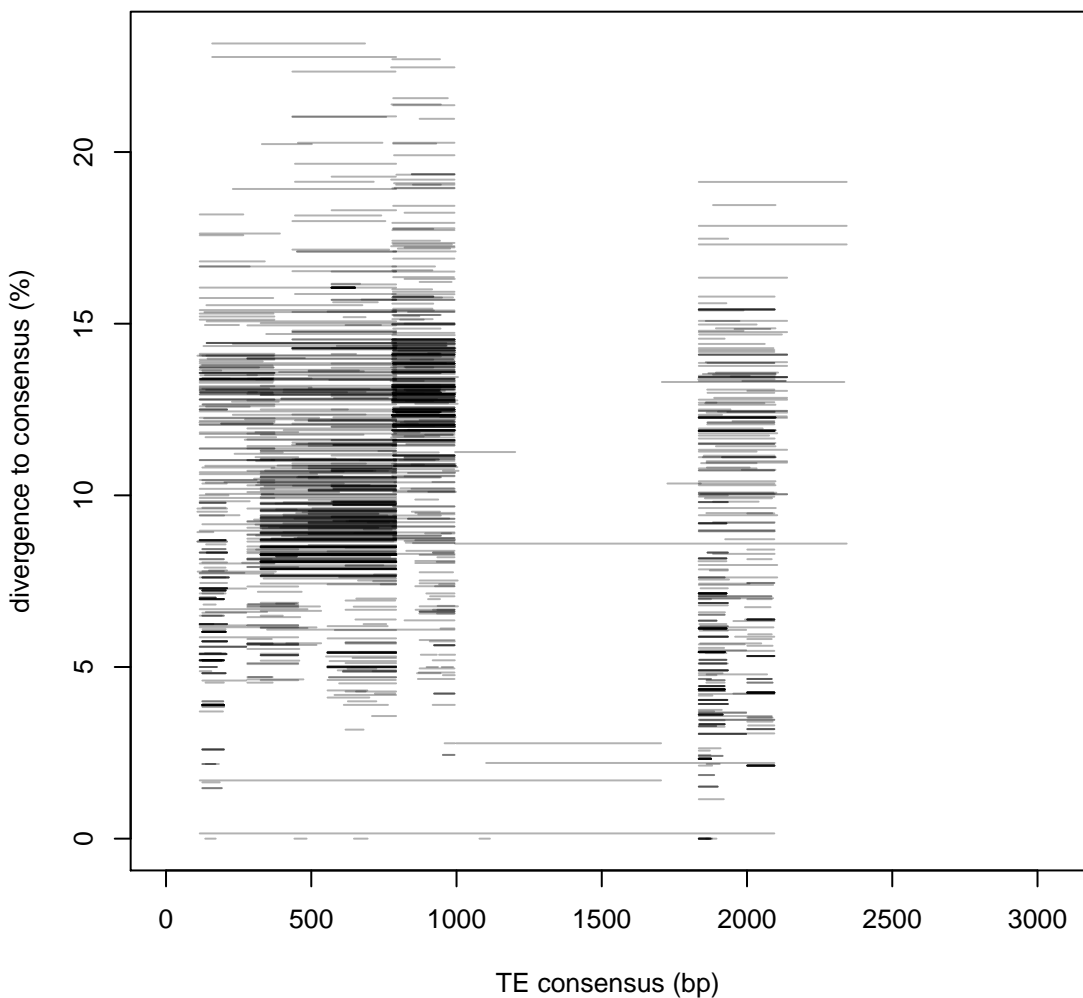
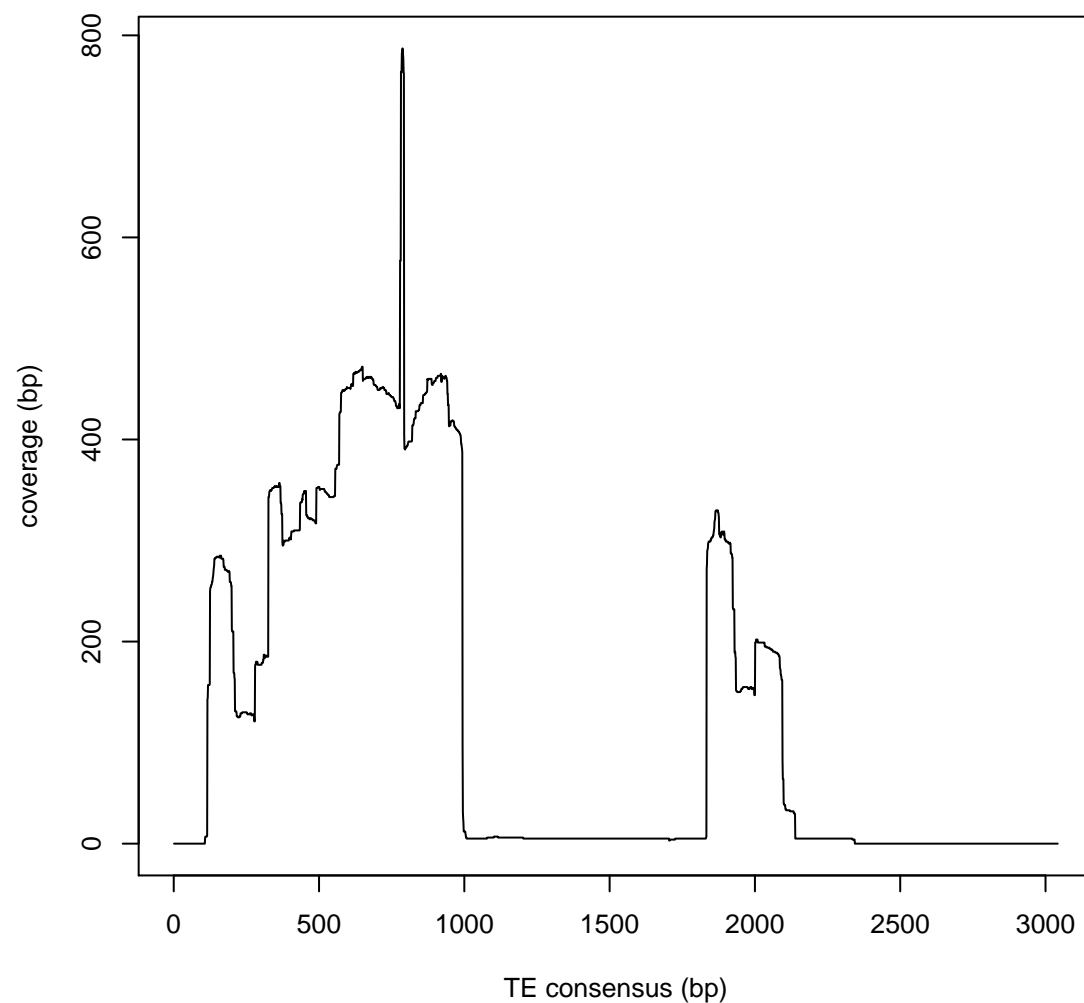


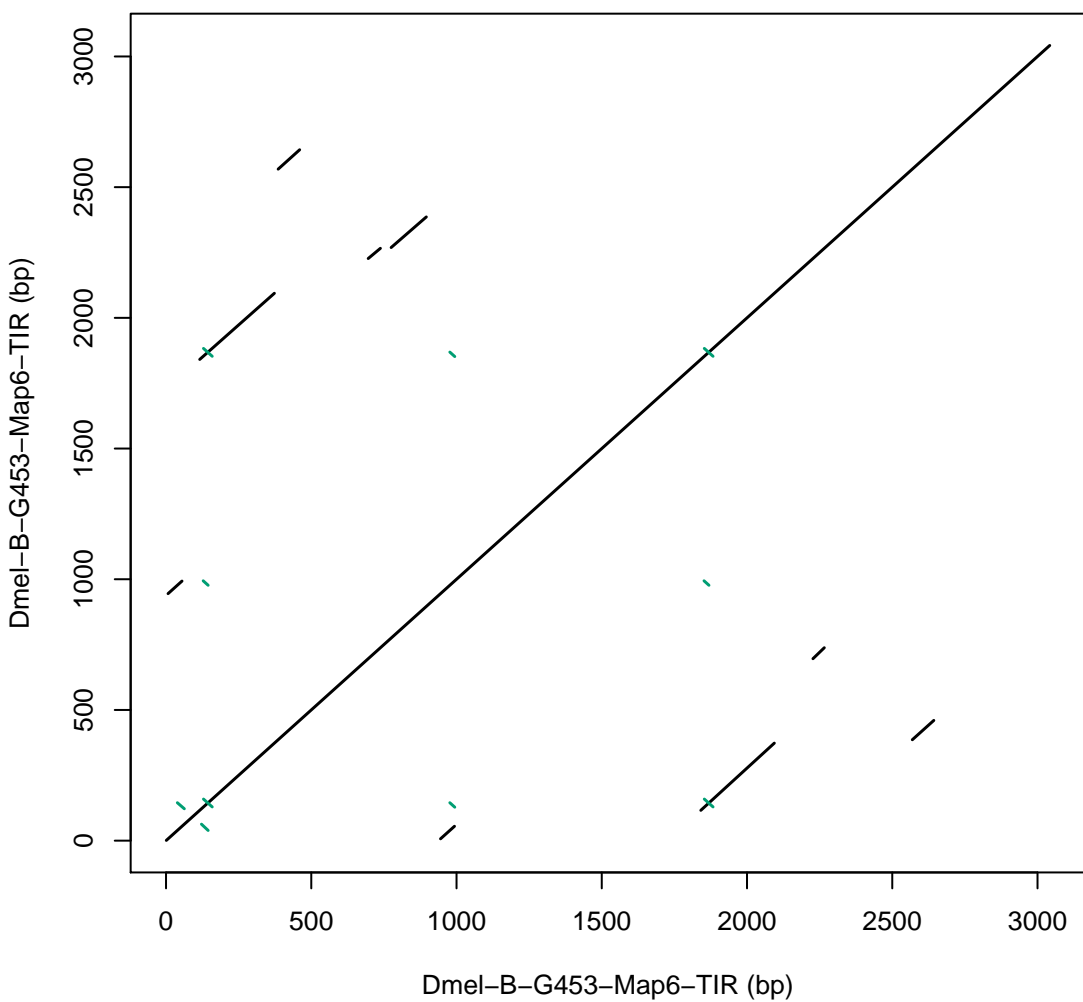
TE: Dmel-B-G453-Map6-TIR
consensus size: 3042bp; fragments: 1712; full length: 0 ($\geq 2737.8\text{bp}$)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

