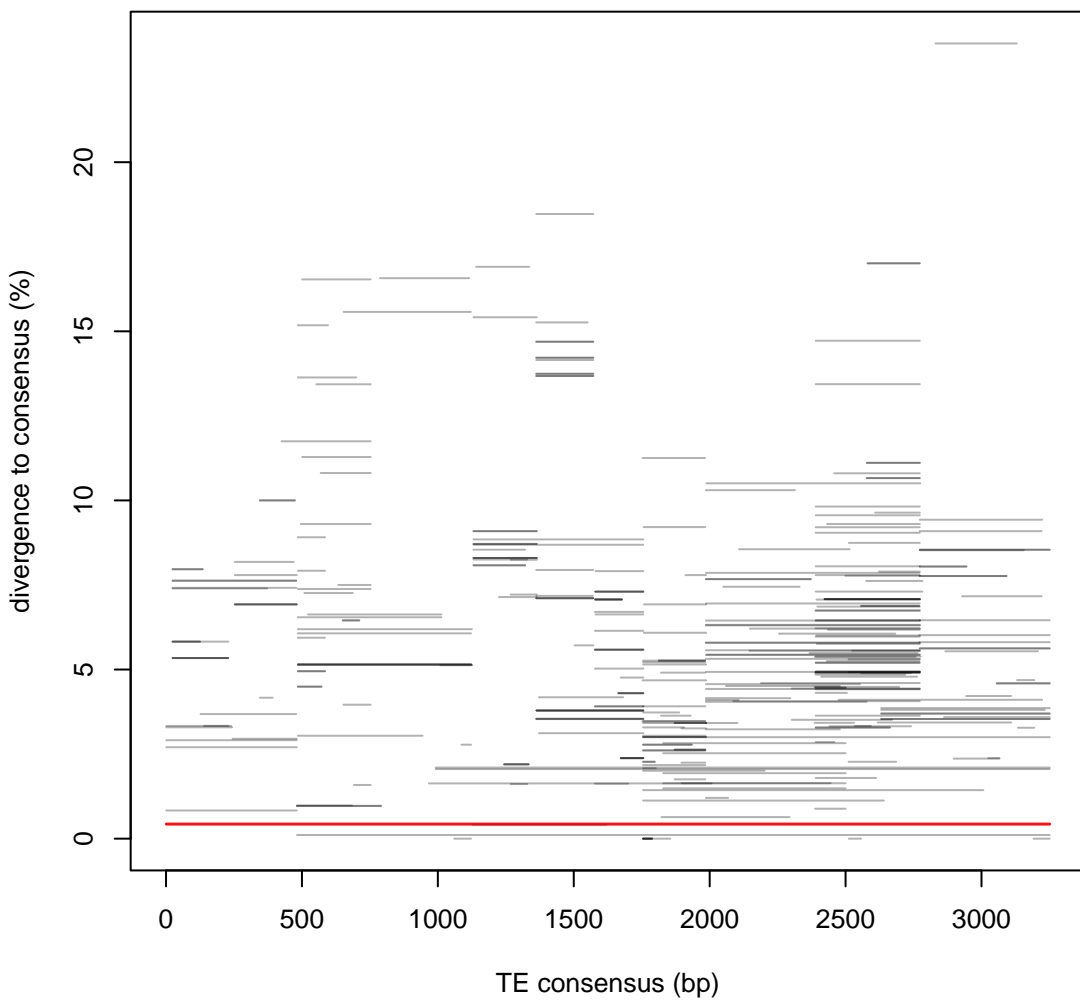
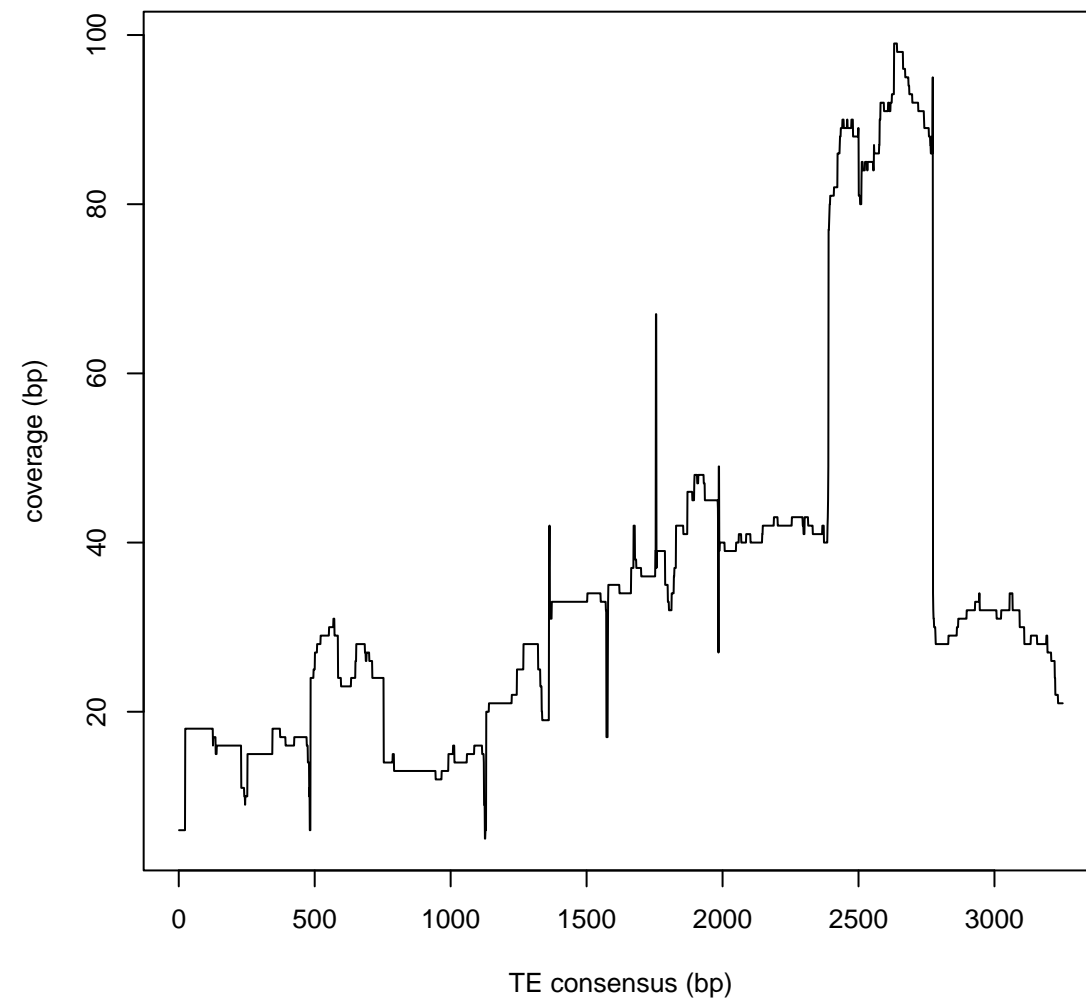


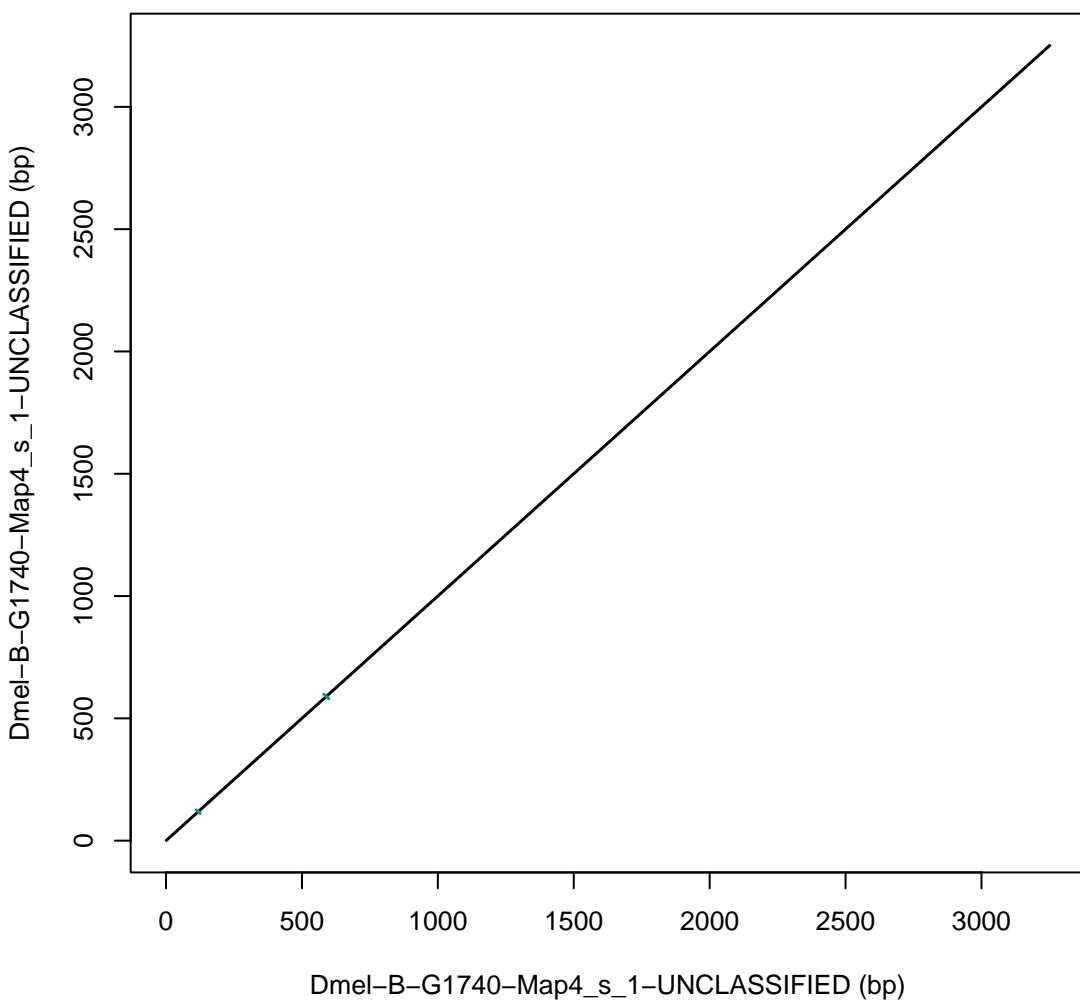
TE: Dmel-B-G1740-Map4\_s\_1-UNCLASSIFIED  
consensus size: 3251bp; fragments: 340; full length: 1 (>=2925.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

