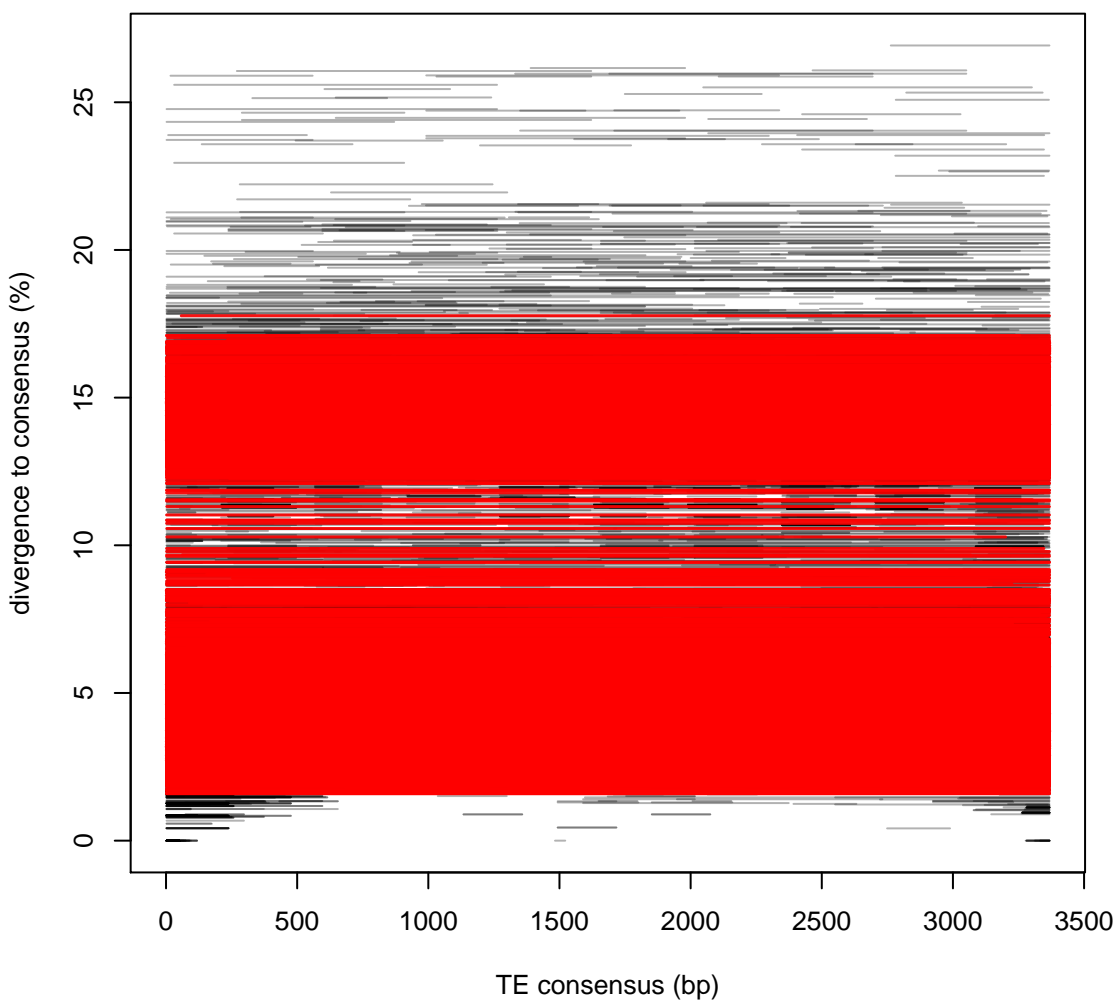
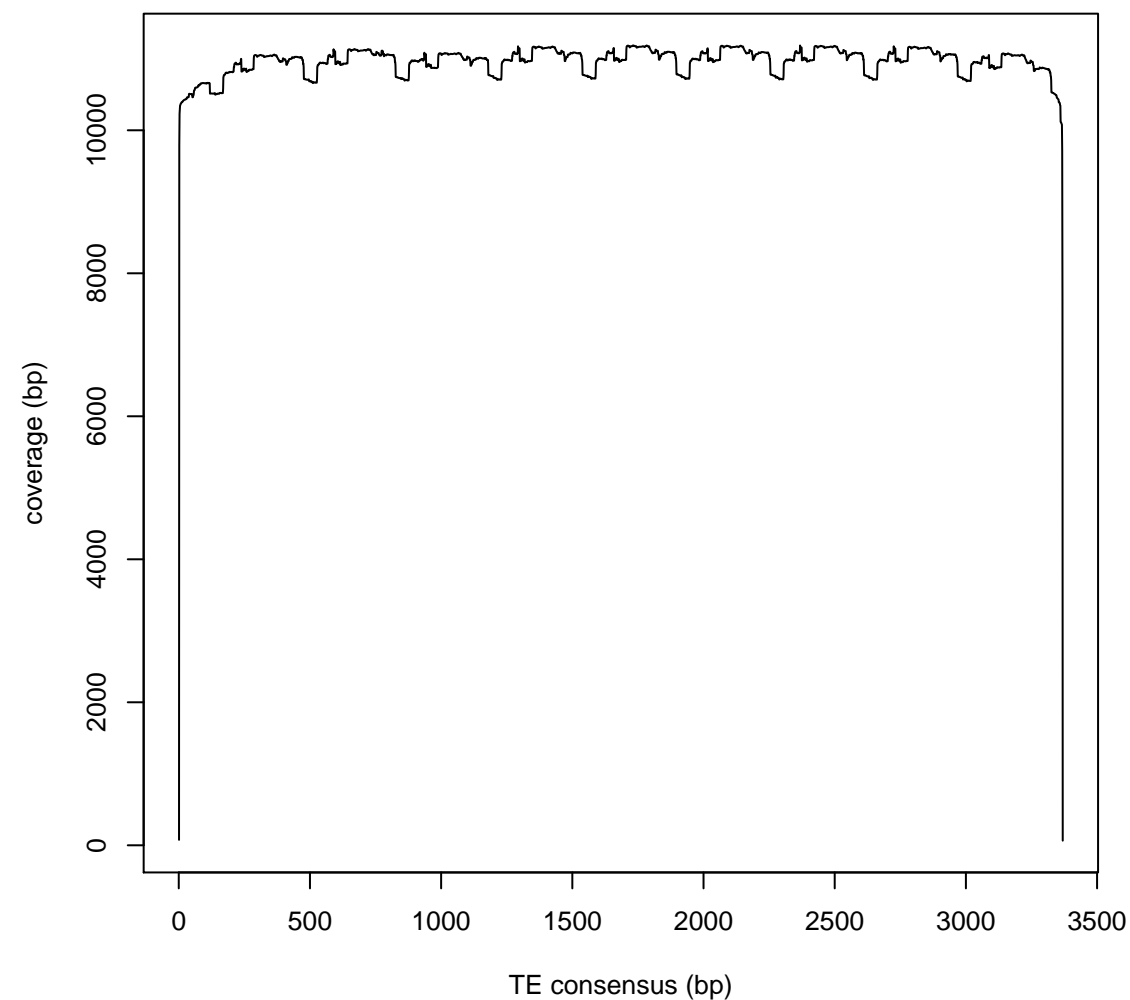


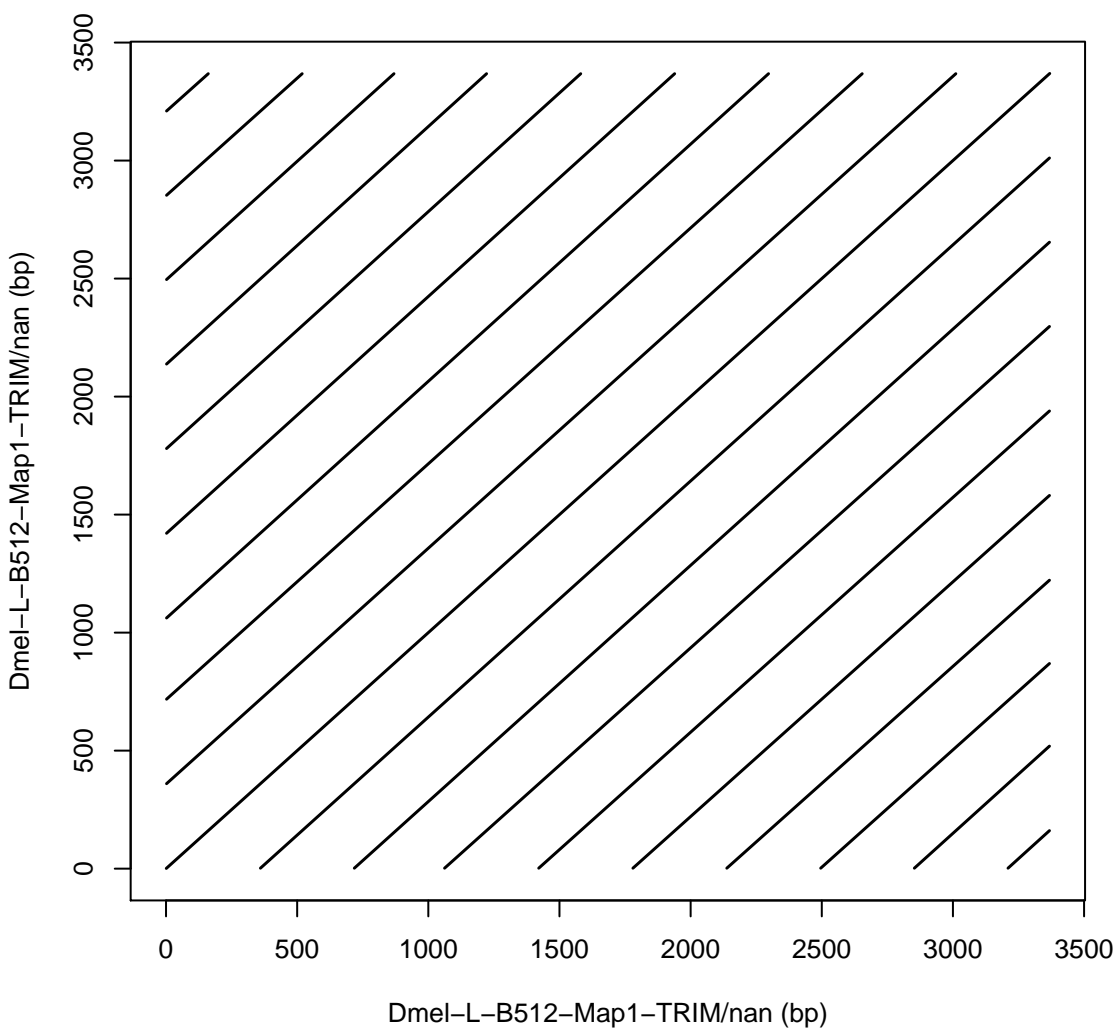
TE: Dmel-L-B512-Map1-TRIM/nan  
consensus size: 3369bp; fragments: 23408; full length: 5714 ( $\geq 3032.1$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

