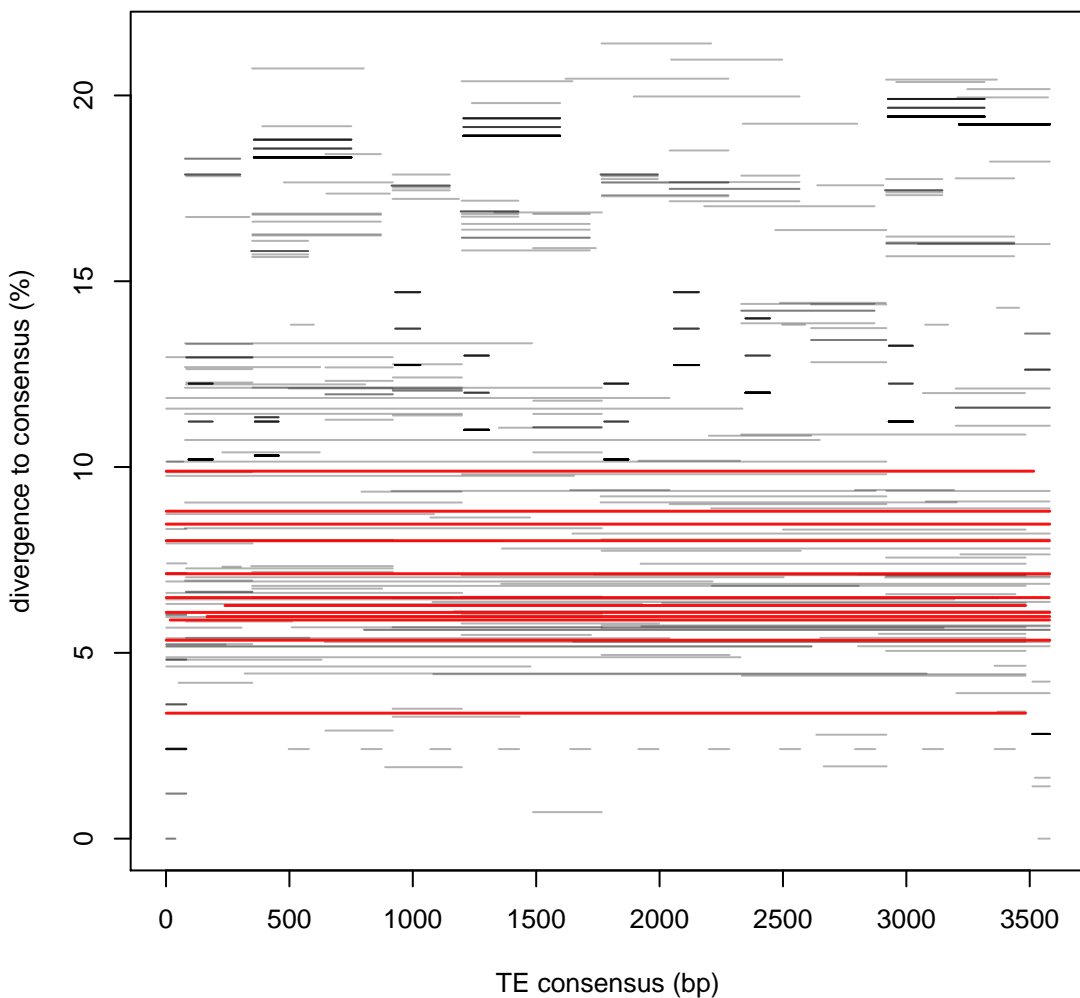
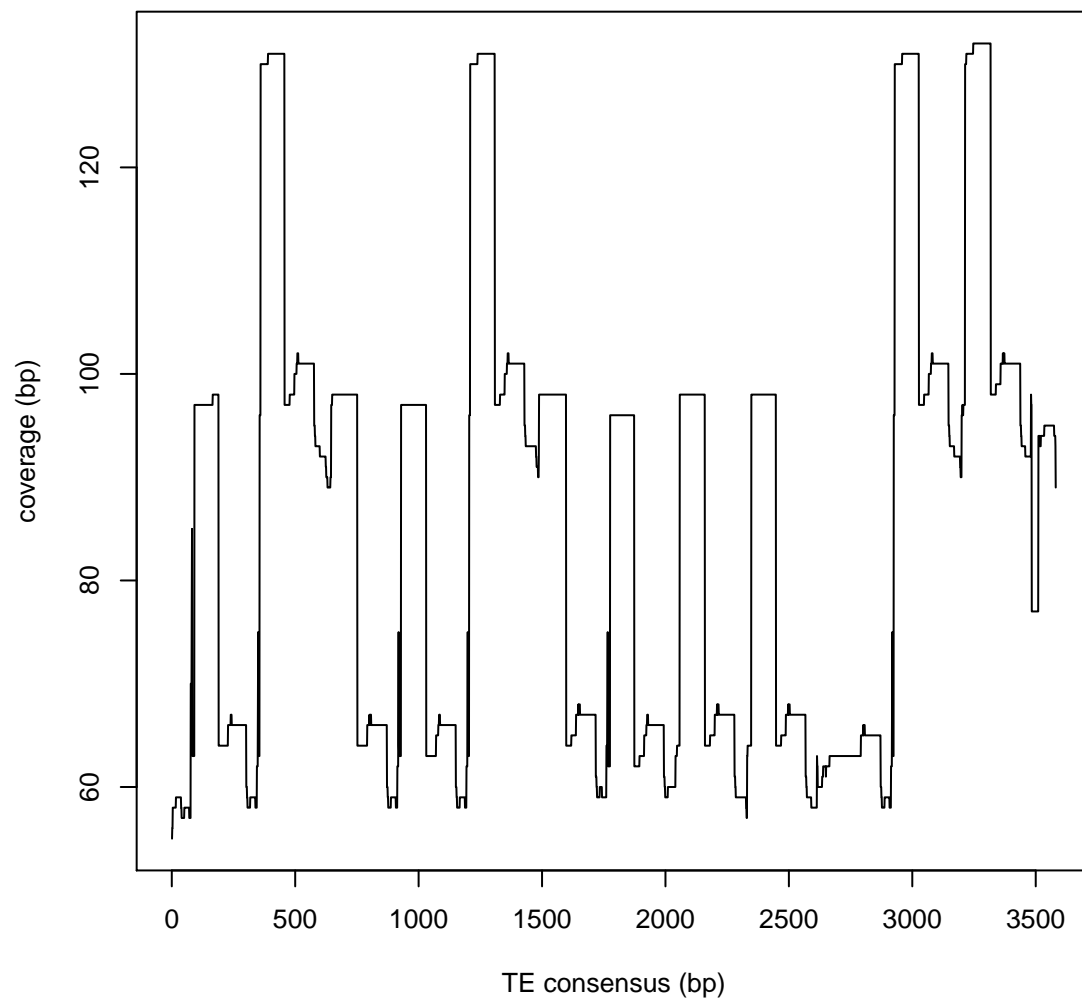


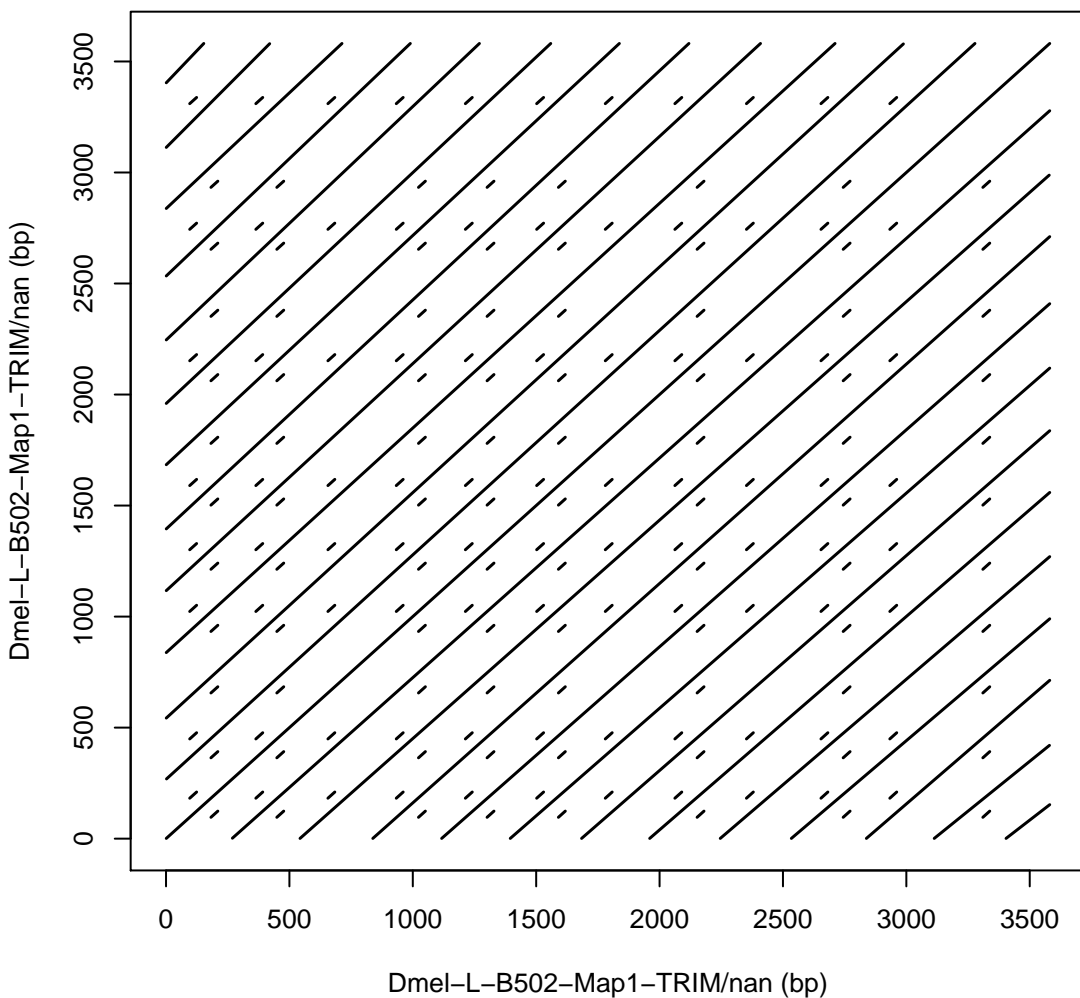
TE: Dmel-L-B502-Map1-TRIM/nan  
consensus size: 3581bp; fragments: 722; full length: 12 ( $\geq 3222.9$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

