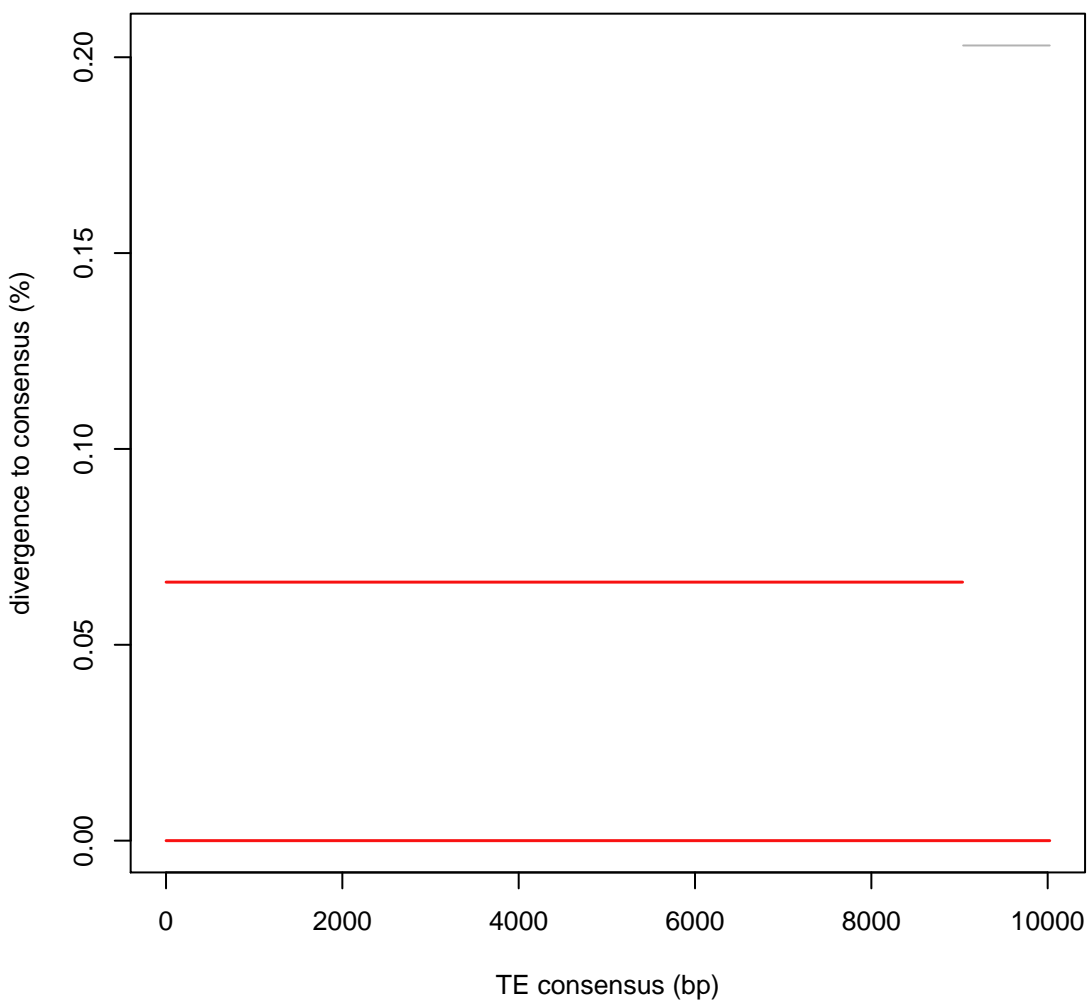
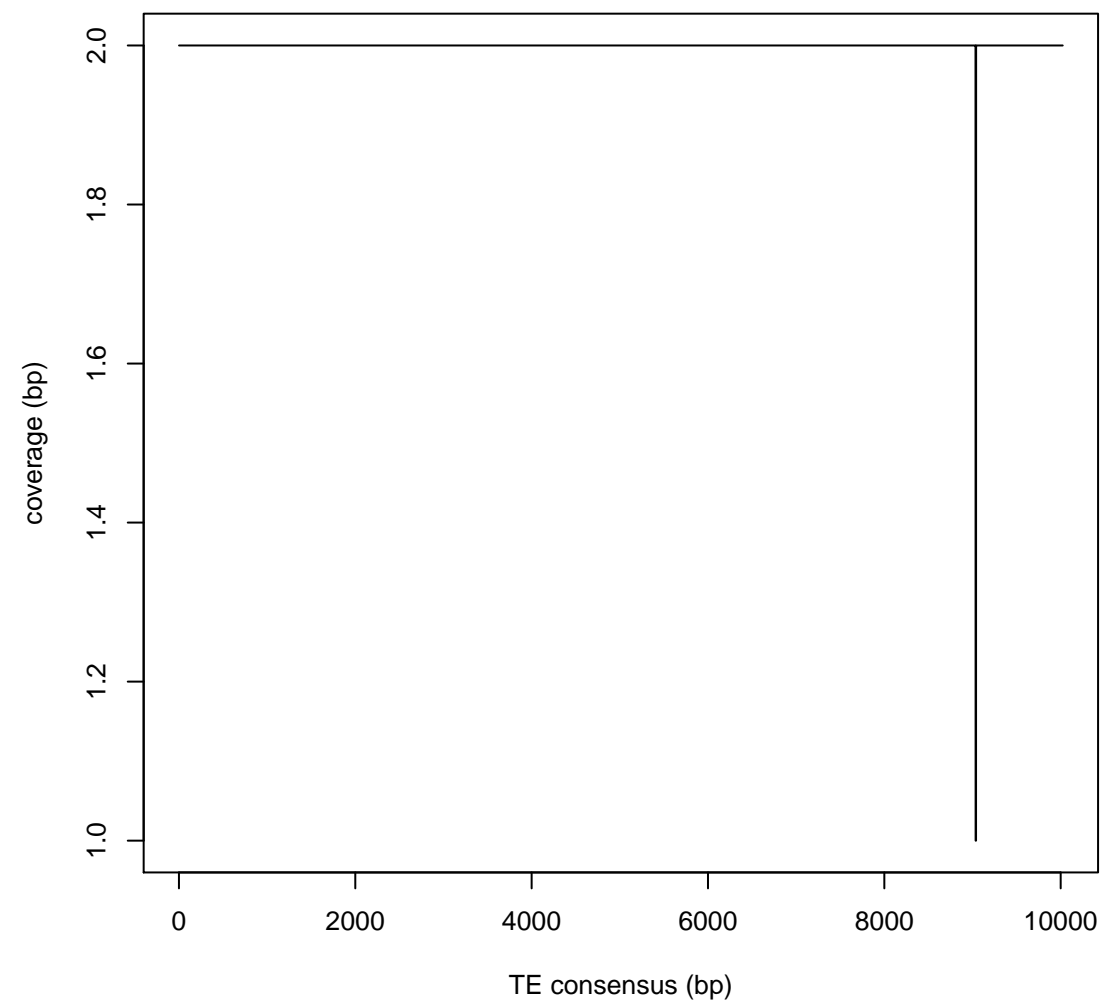


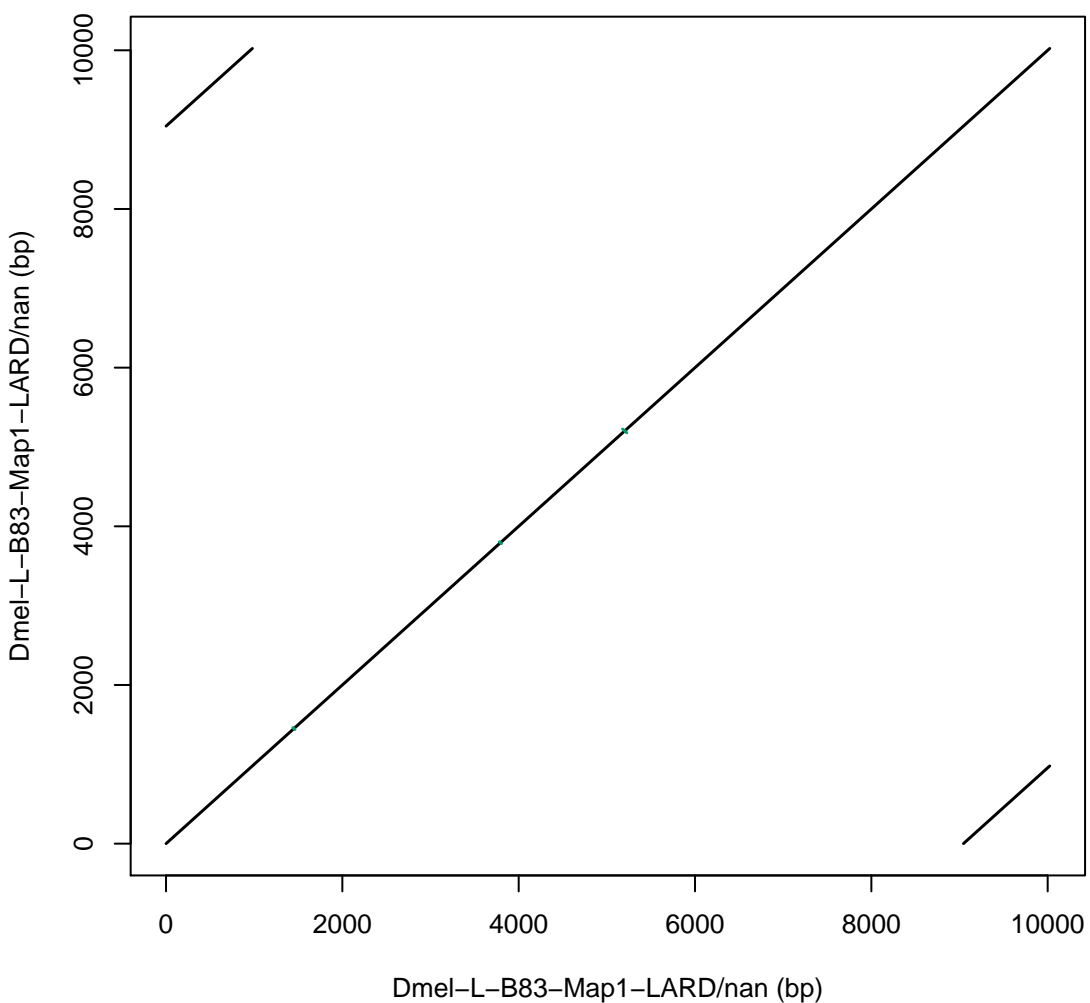
TE: Dmel-L-B83-Map1-LARD/nan  
consensus size: 10023bp; fragments: 3; full length: 2 ( $\geq 9020.7$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

