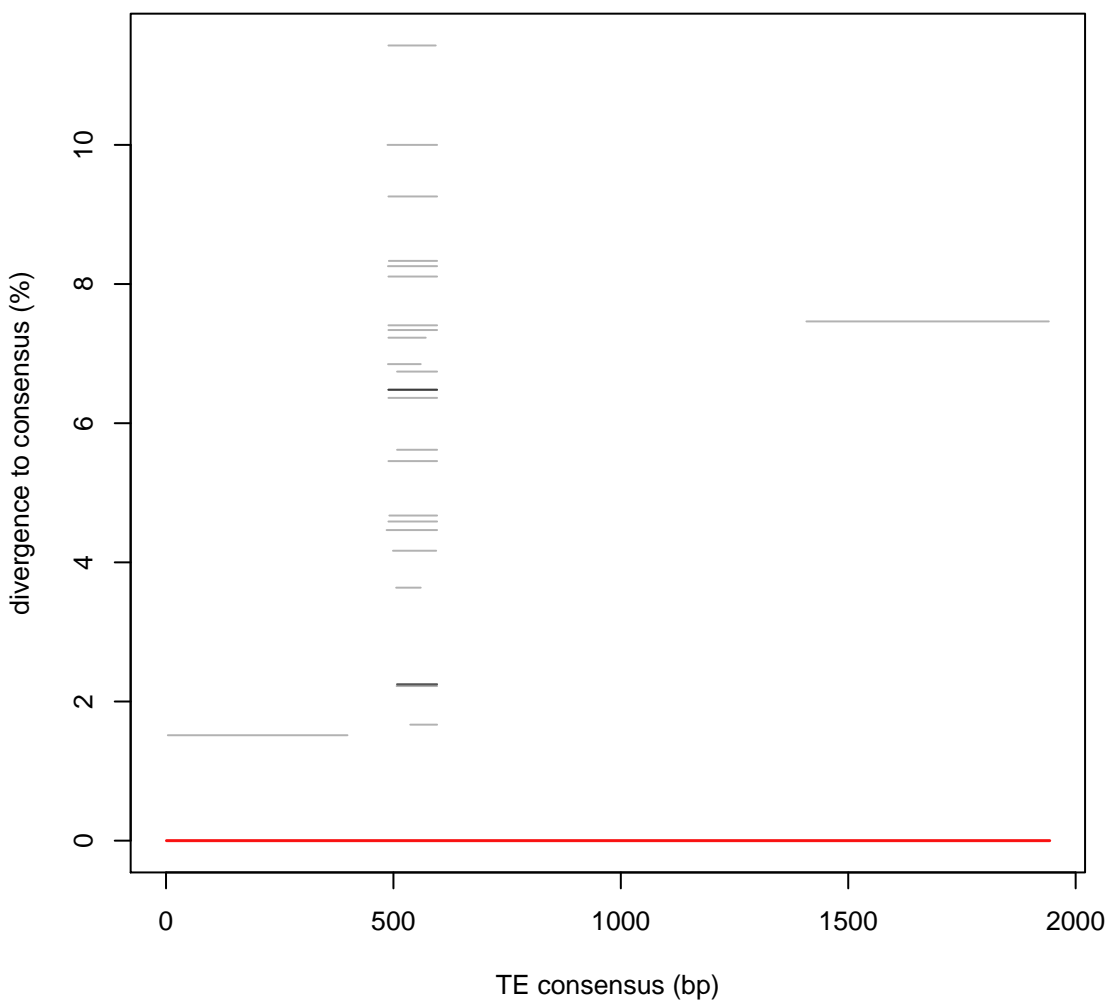
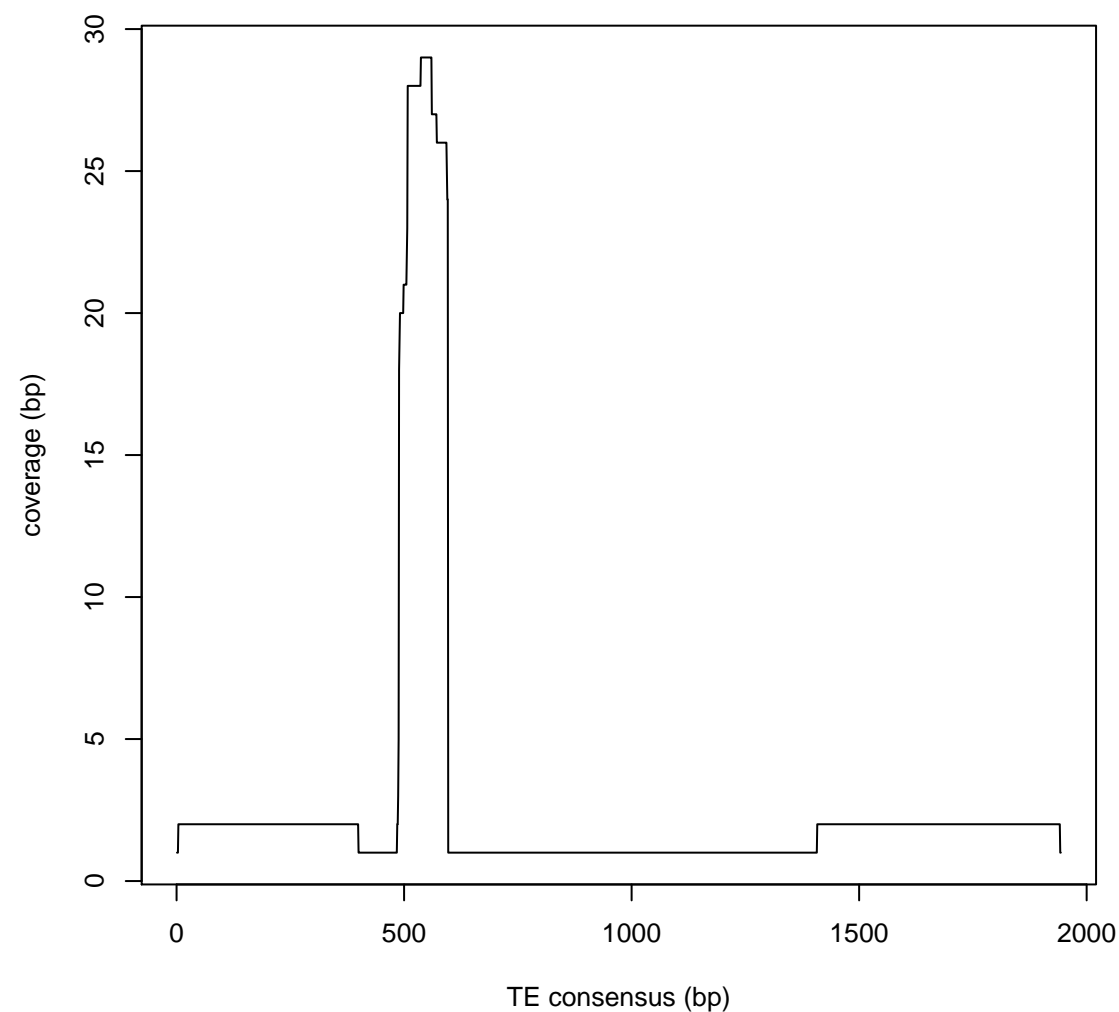


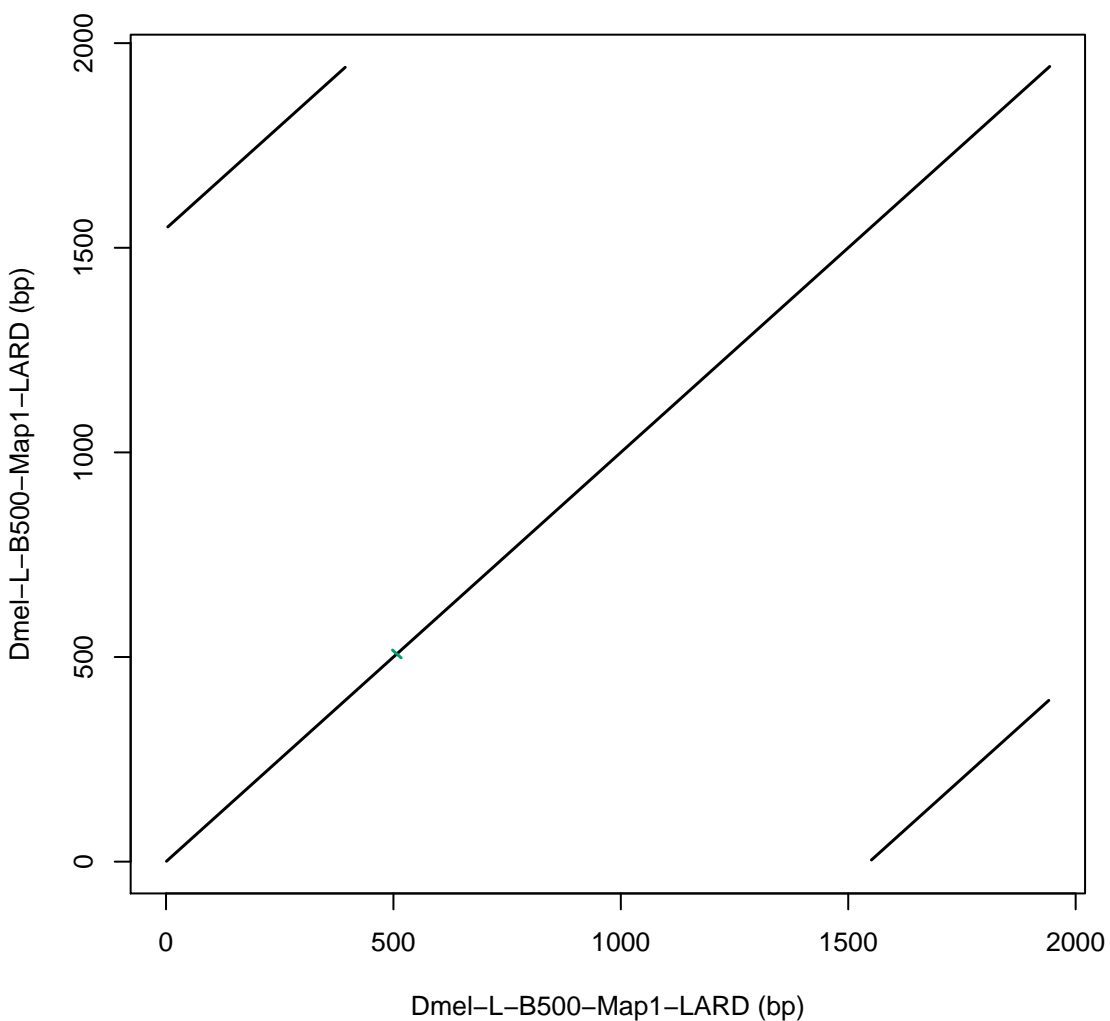
TE: Dmel-L-B500-Map1-LARD  
consensus size: 1943bp; fragments: 31; full length: 1 ( $\geq 1748.7$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

