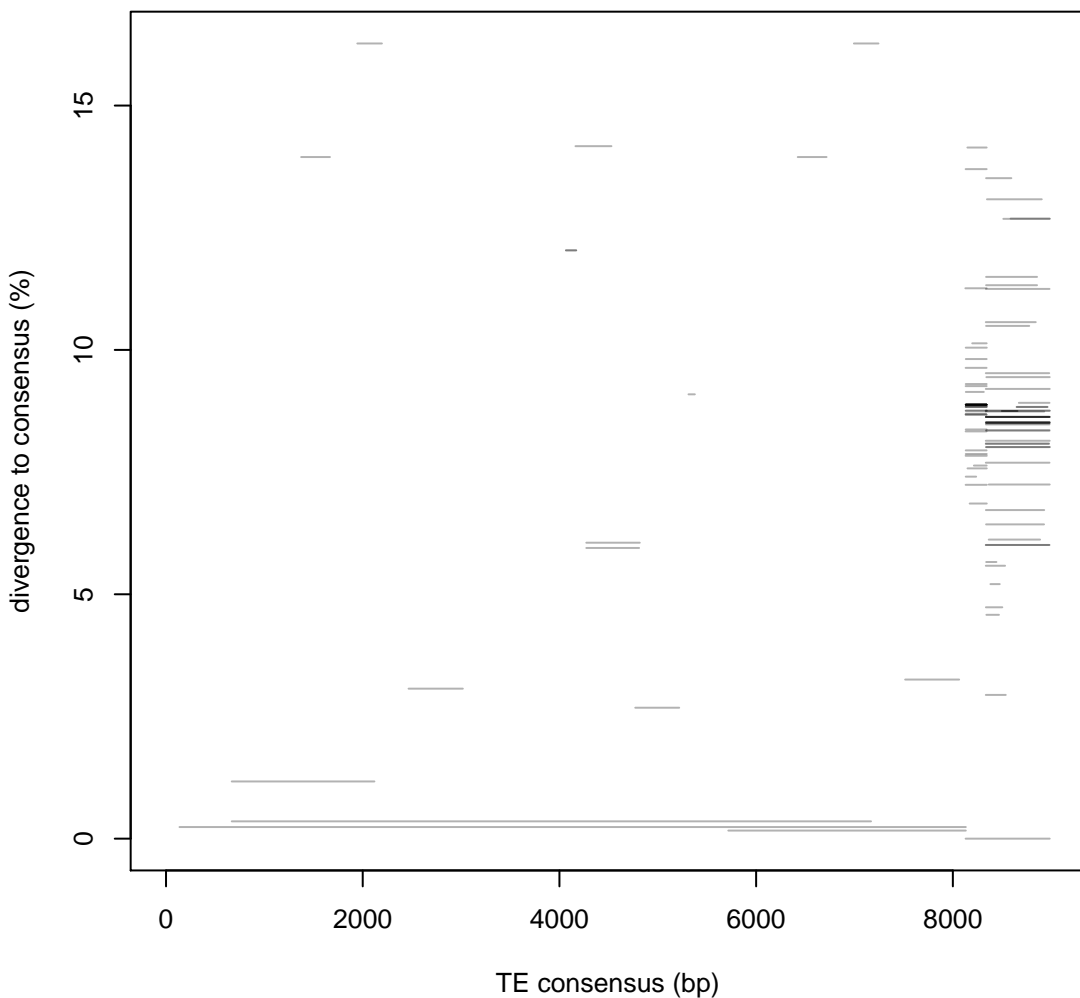
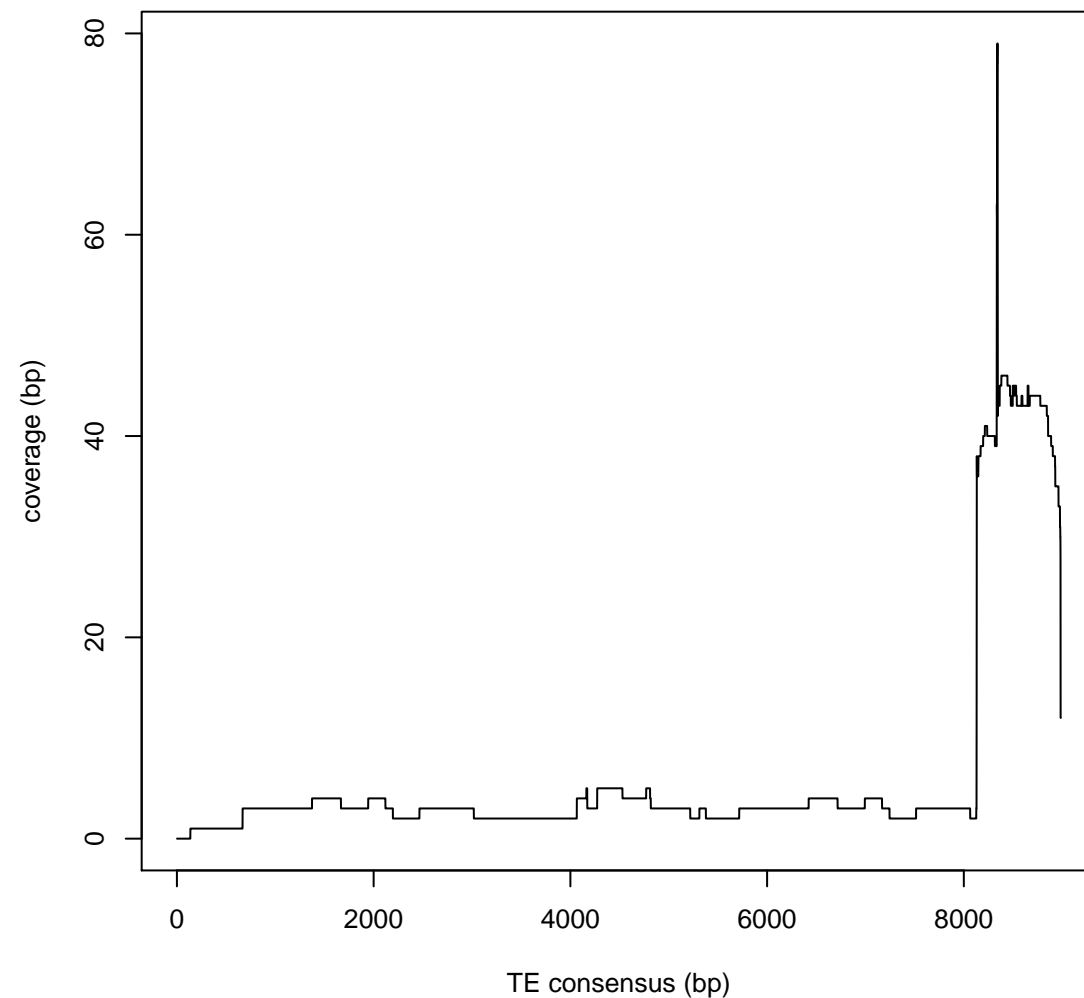


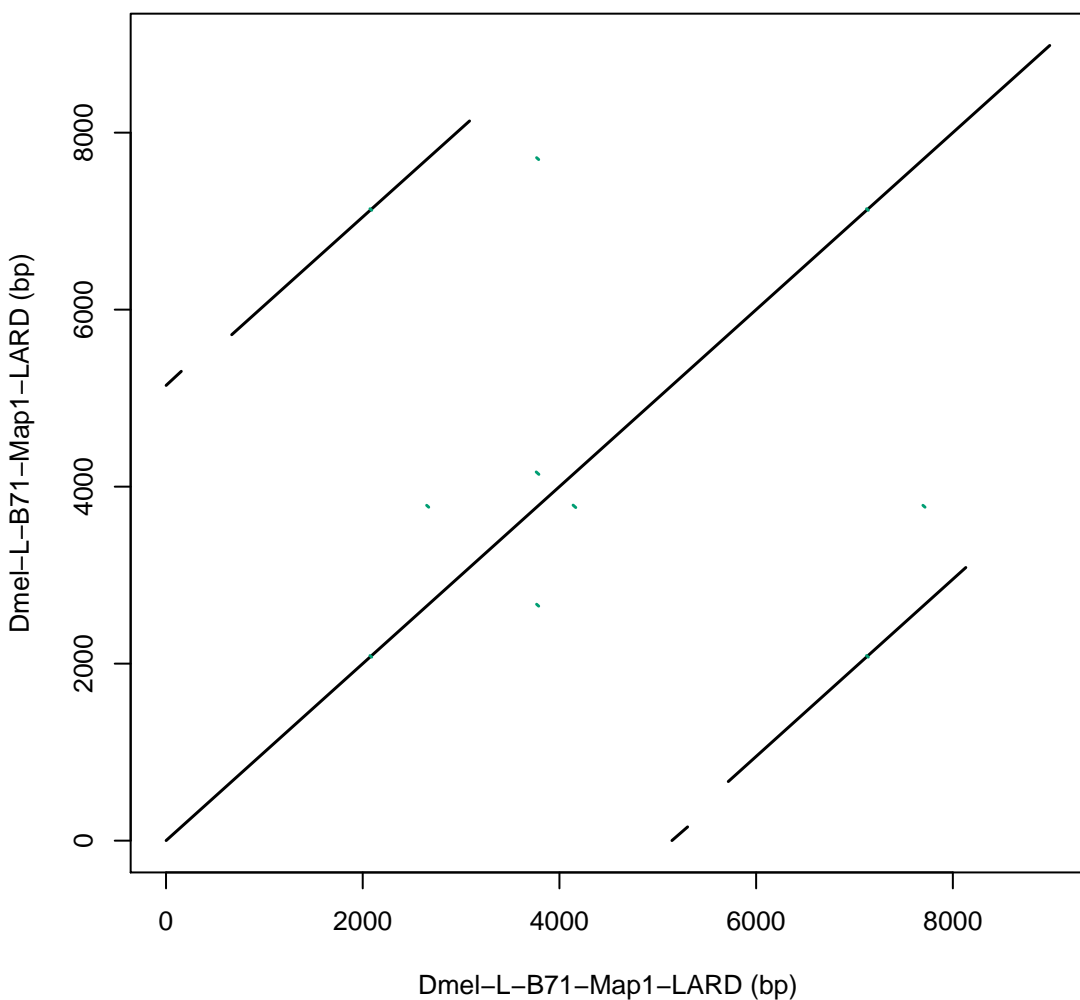
TE: Dmel-L-B71-Map1-LARD  
consensus size: 8985bp; fragments: 110; full length: 0 (>=8086.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

