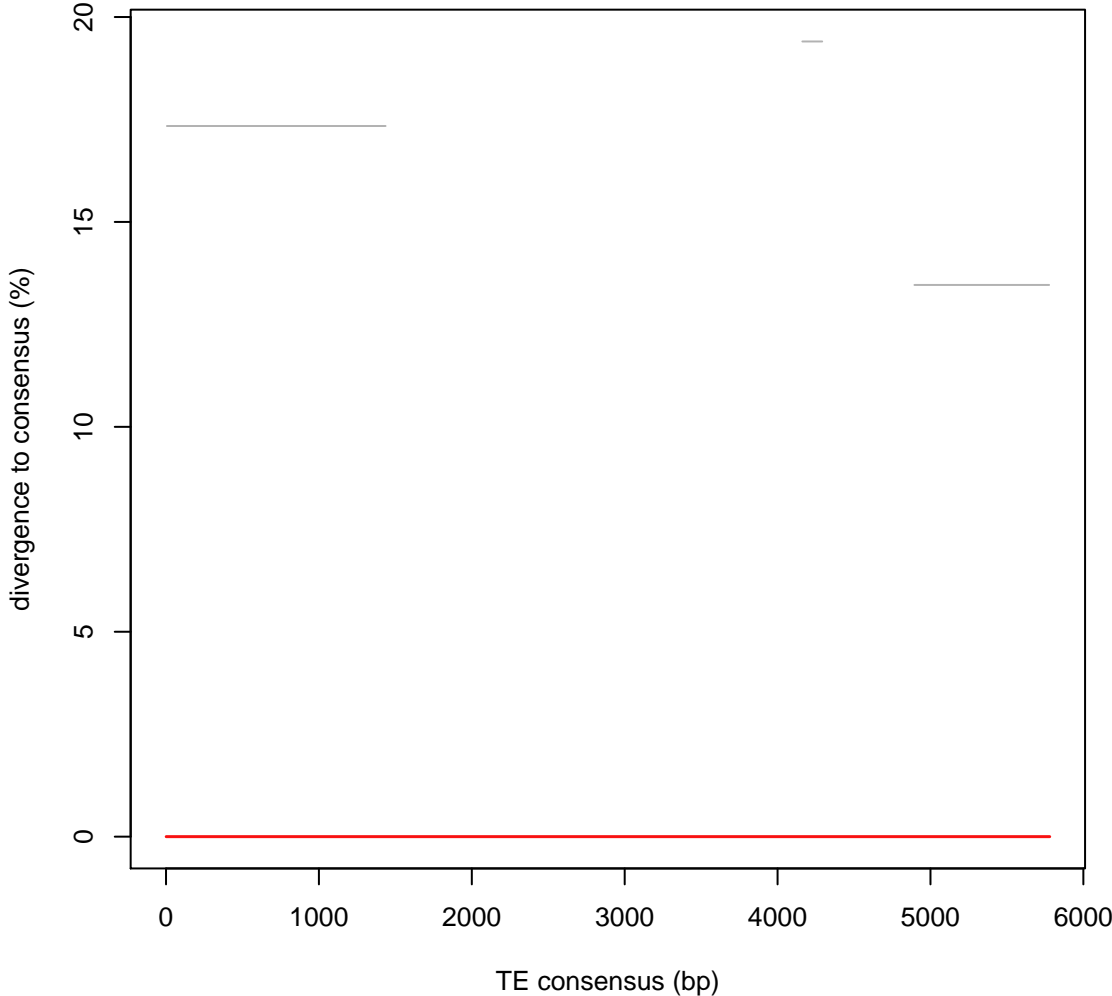
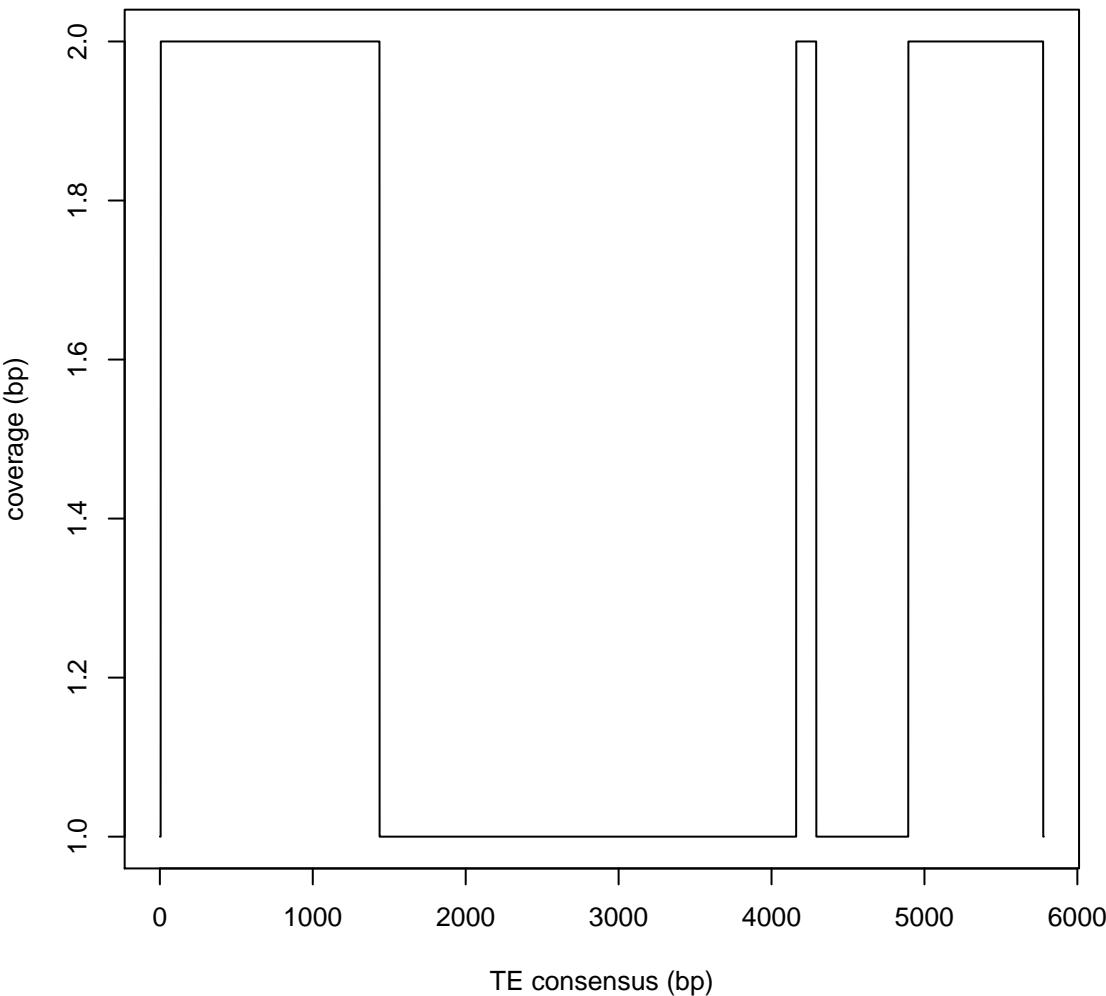


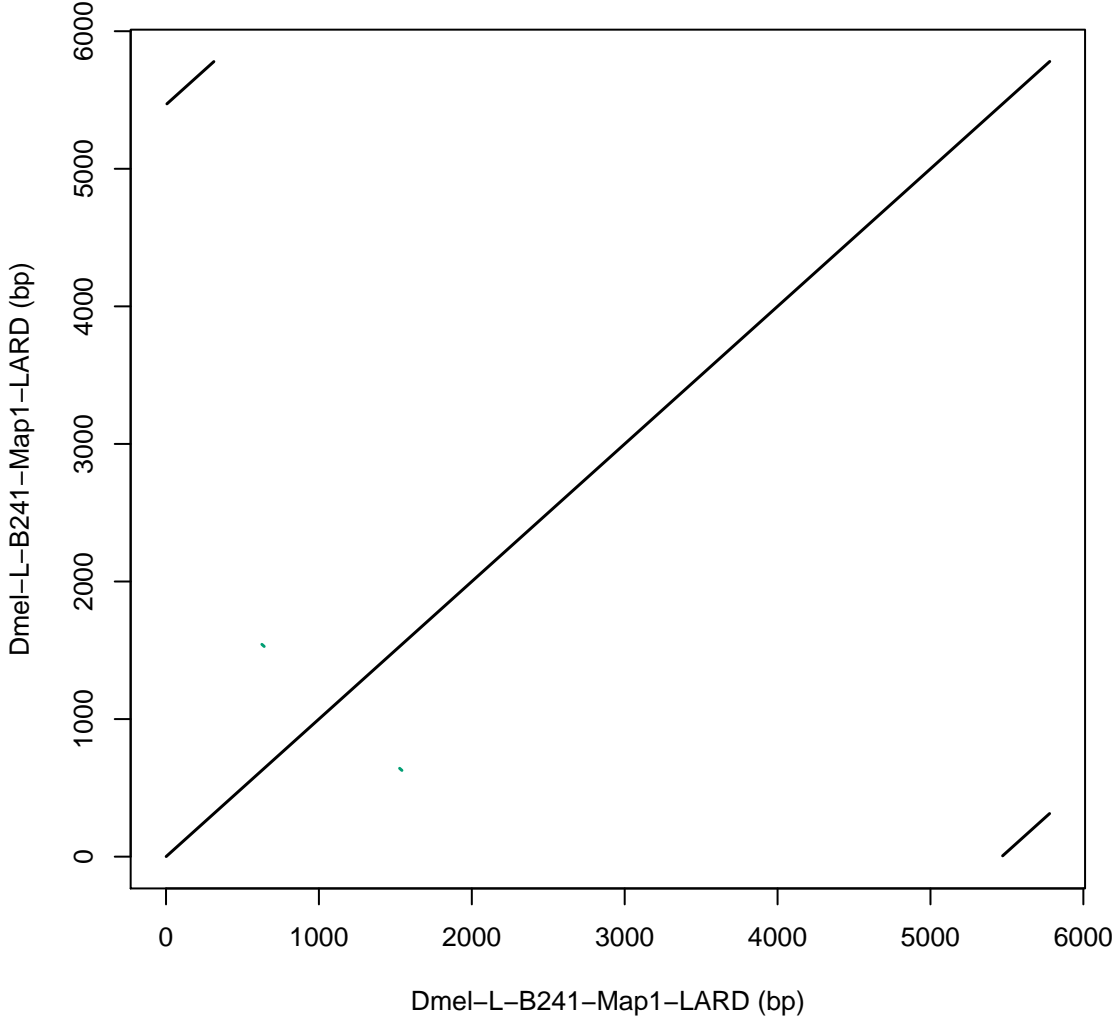
TE: Dmel-L-B241-Map1-LARD  
consensus size: 5780bp; fragments: 4; full length: 1 (>=5202bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

