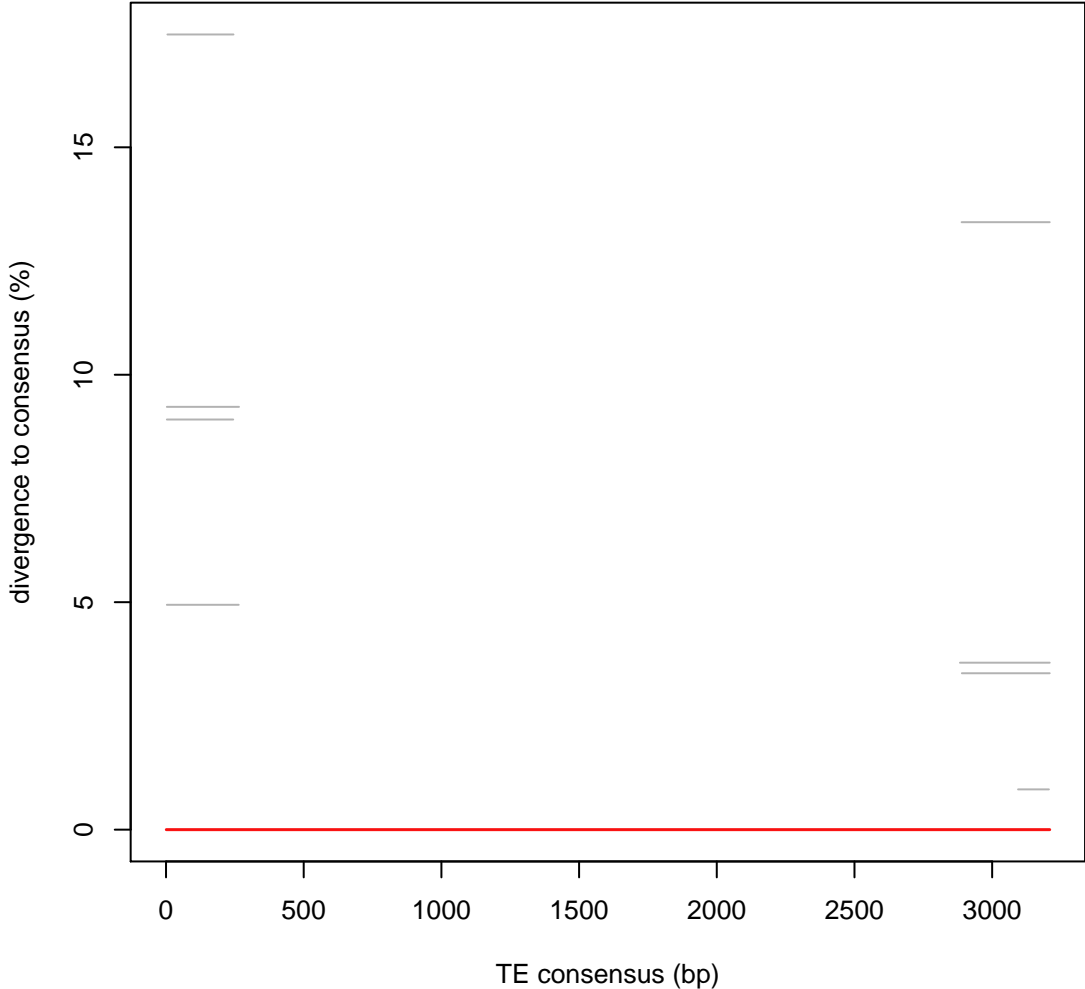
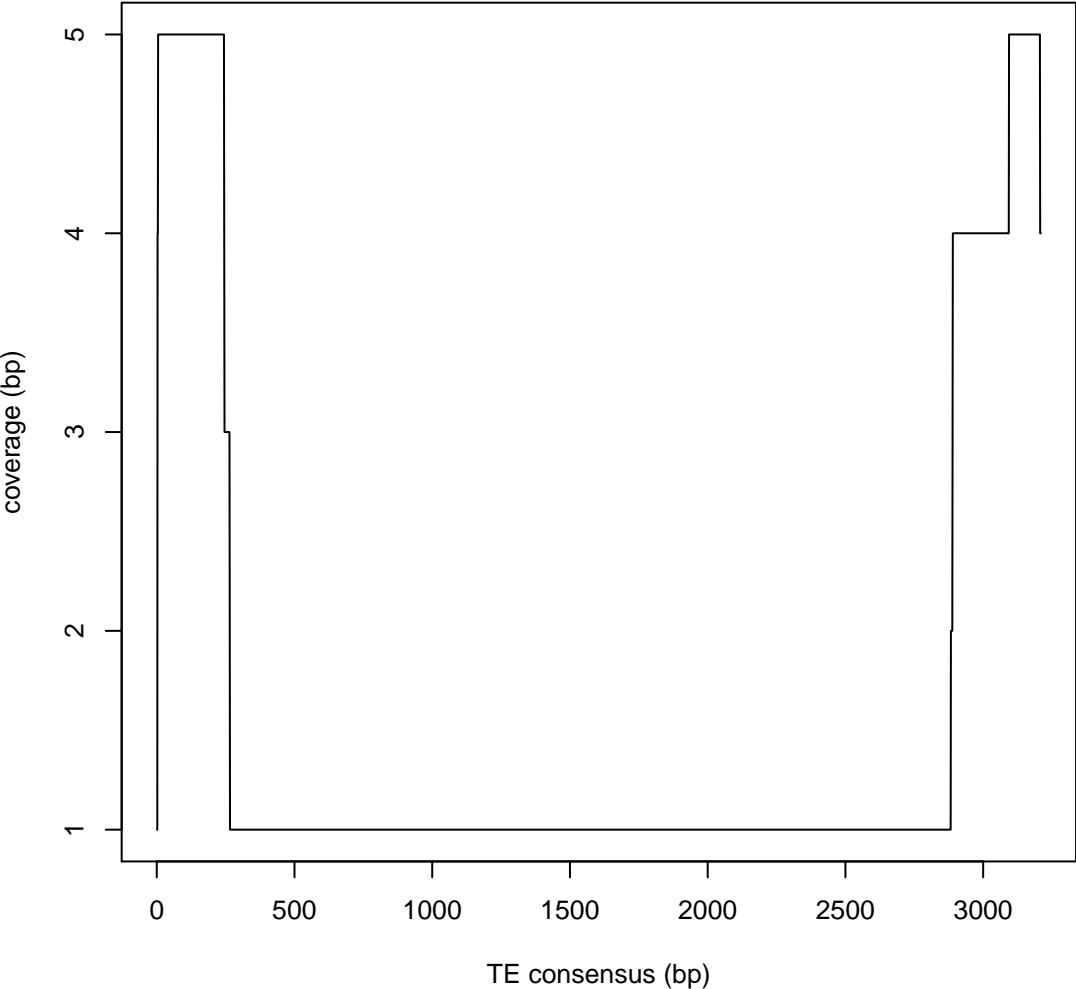


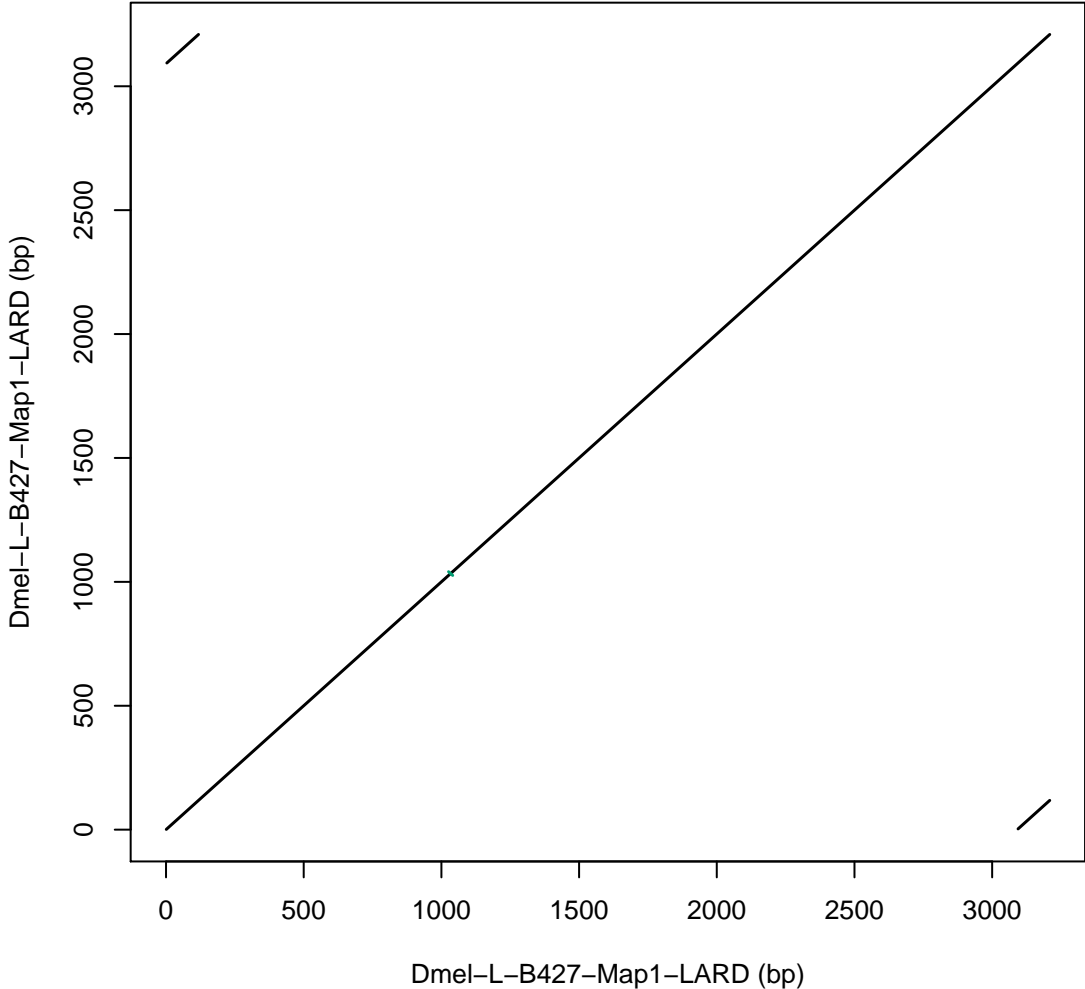
TE: Dmel-L-B427-Map1-LARD  
consensus size: 3209bp; fragments: 9; full length: 1 (>=2888.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

