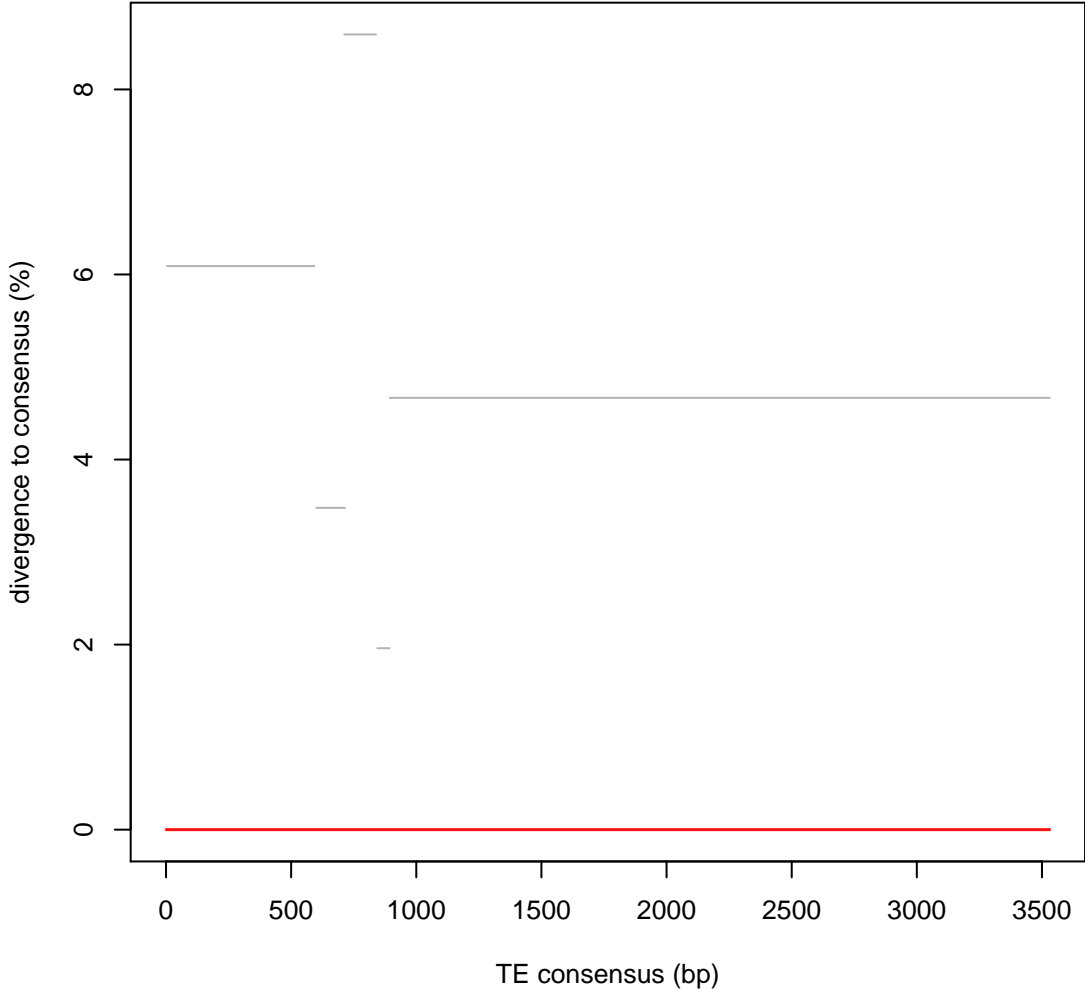
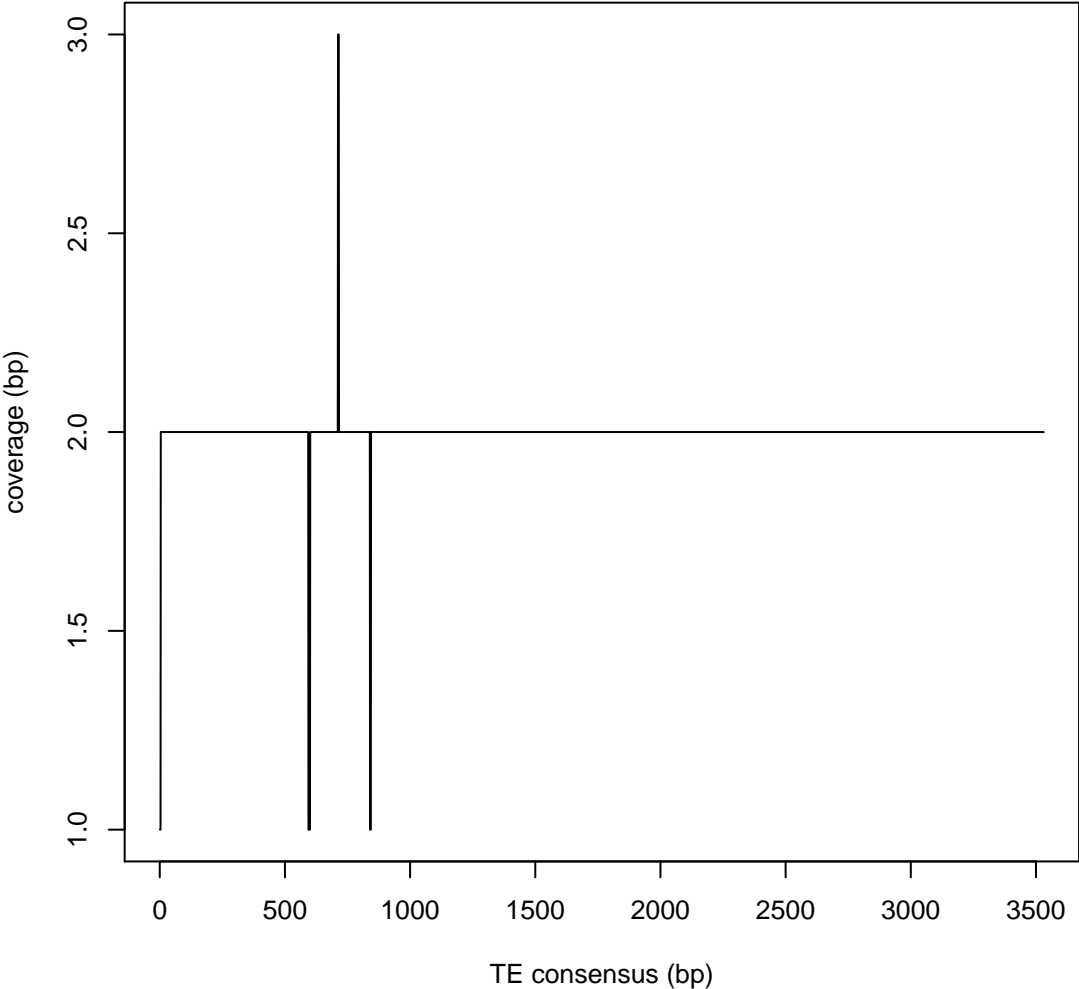


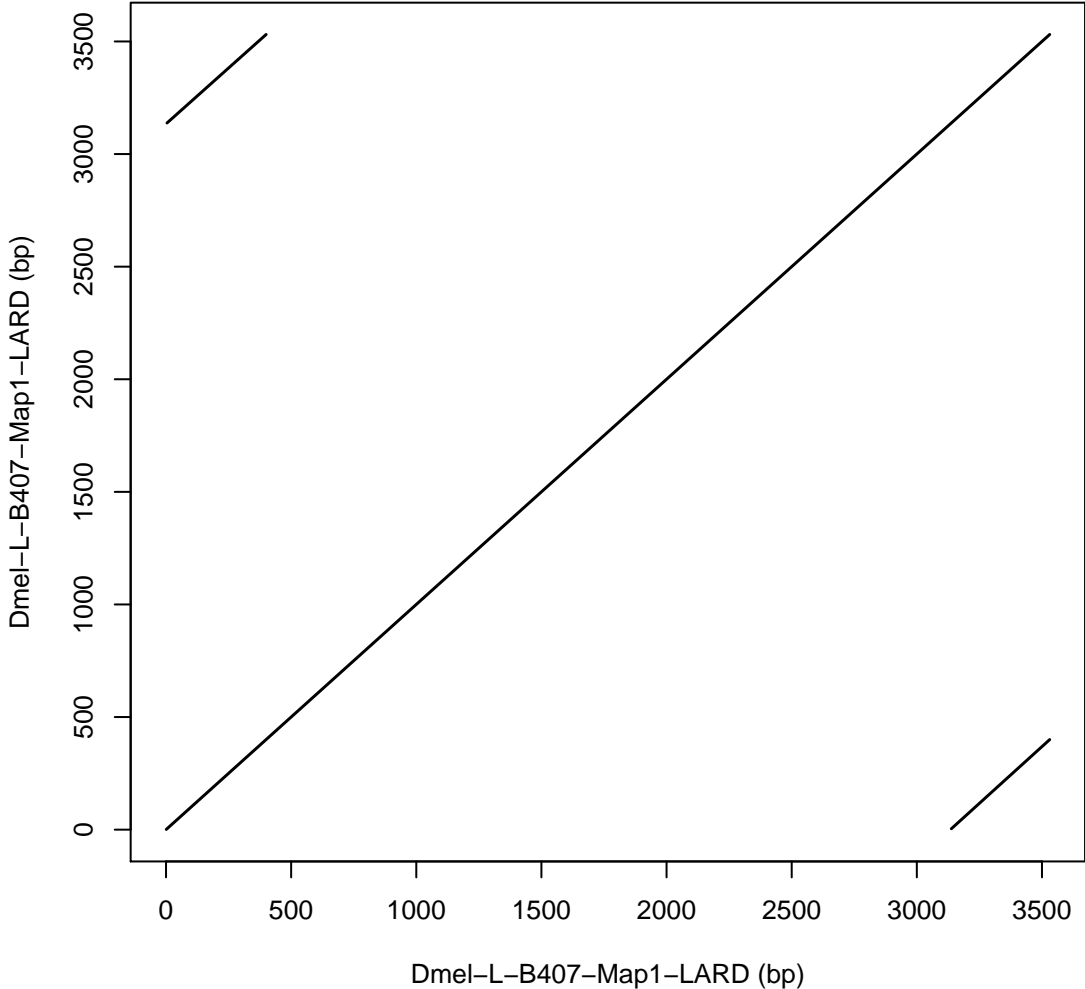
TE: Dmel-L-B407-Map1-LARD  
consensus size: 3531bp; fragments: 6; full length: 1 (>=3177.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

