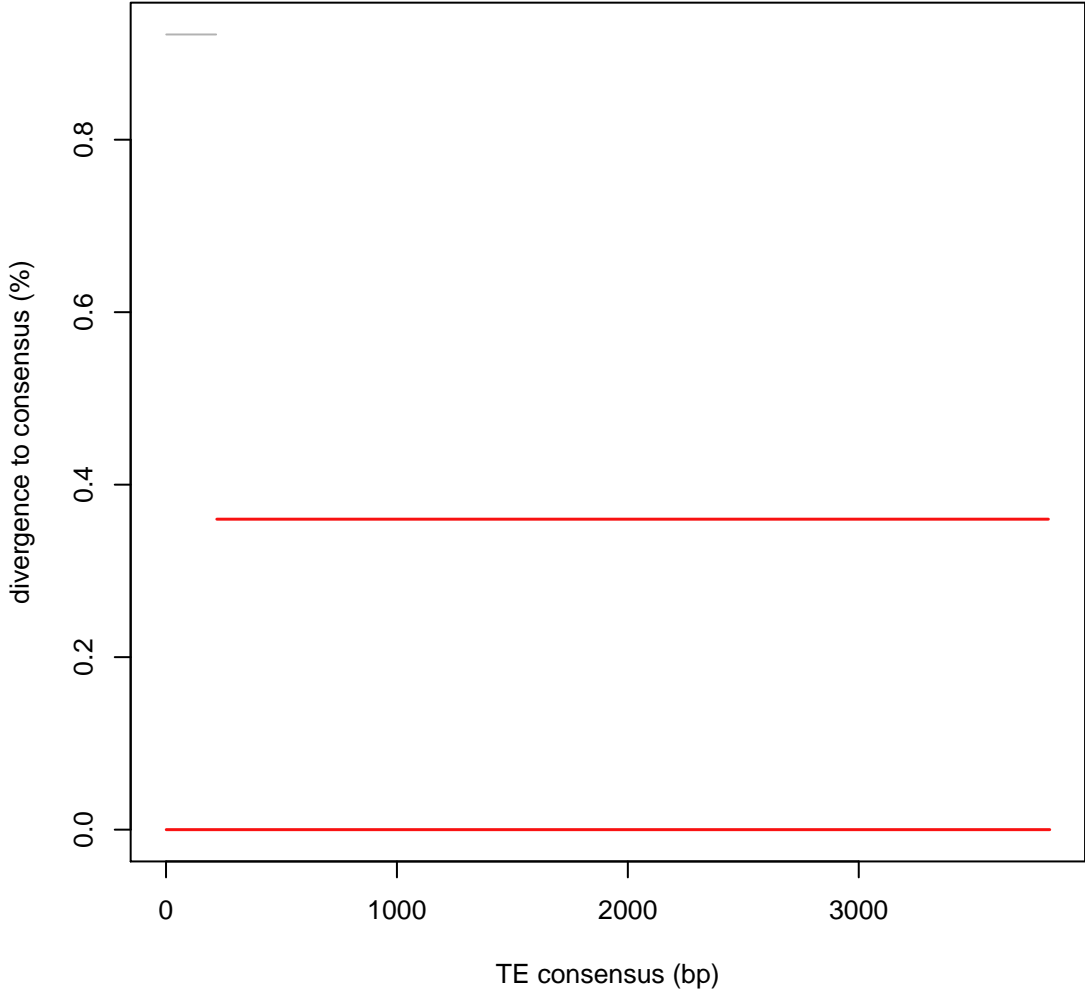
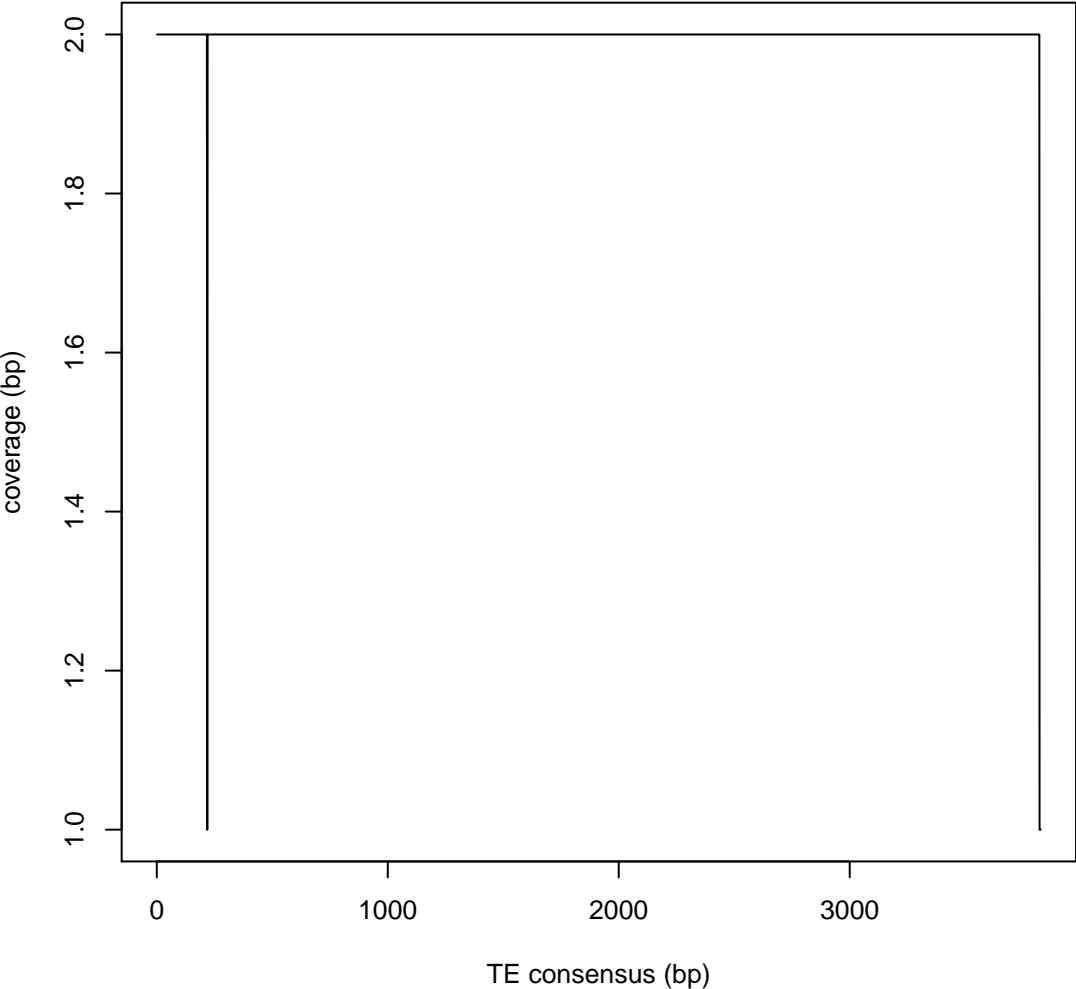


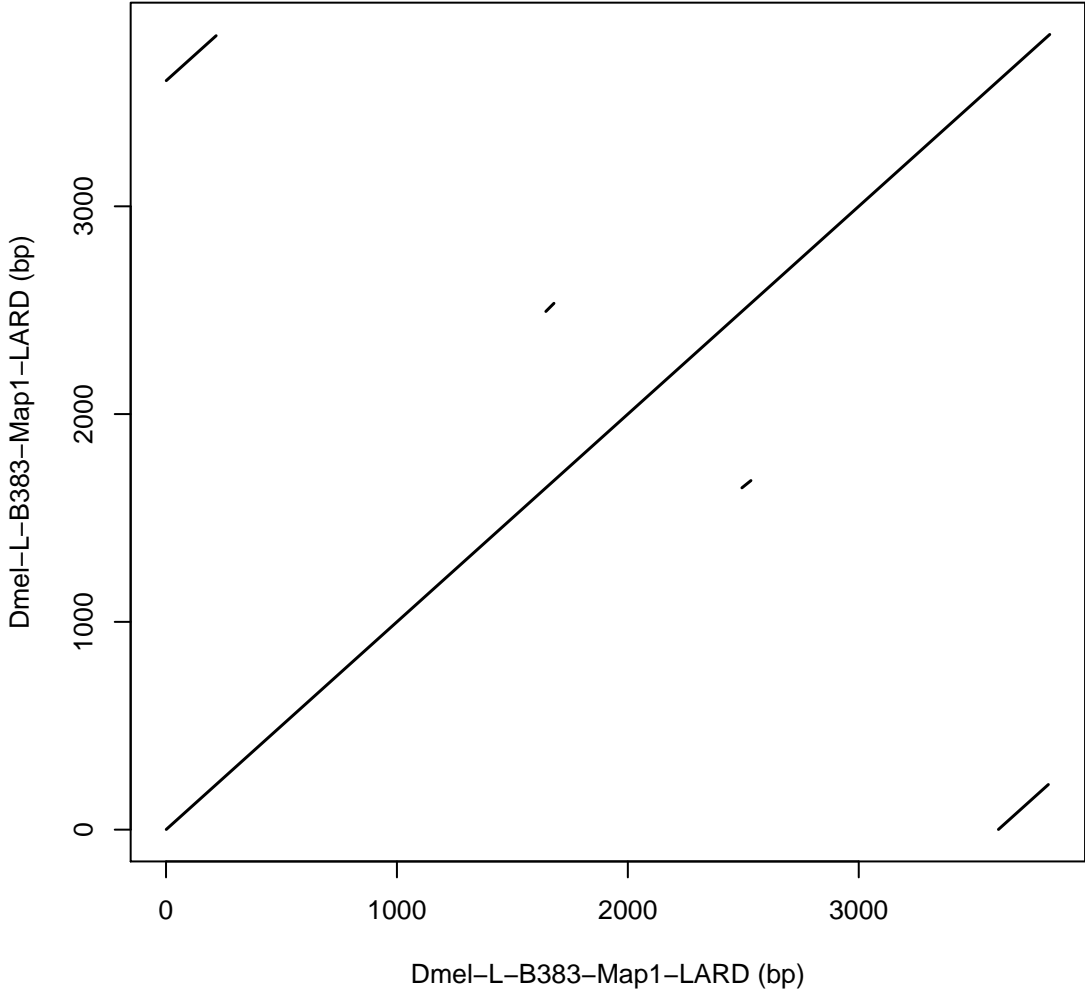
TE: Dmel-L-B383-Map1-LARD  
consensus size: 3827bp; fragments: 3; full length: 2 (>=3444.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

