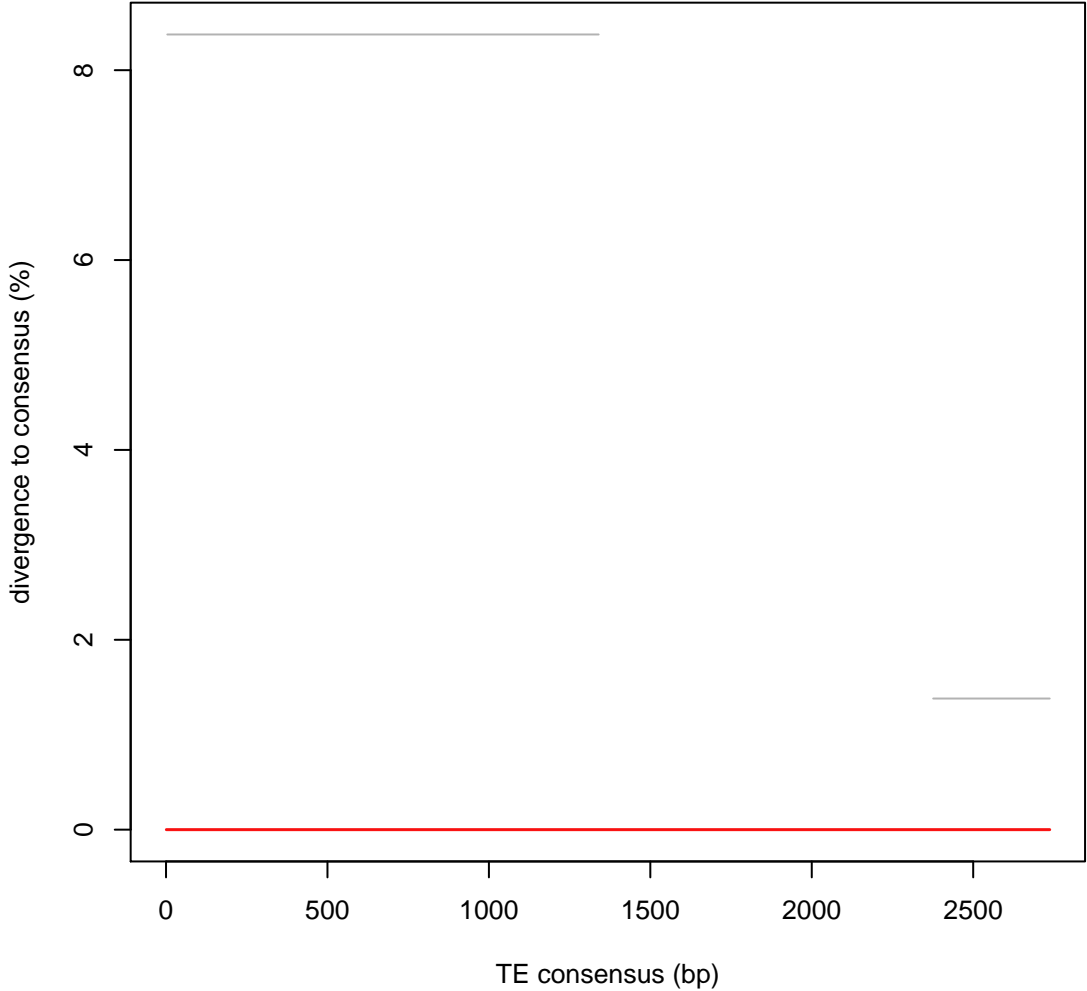
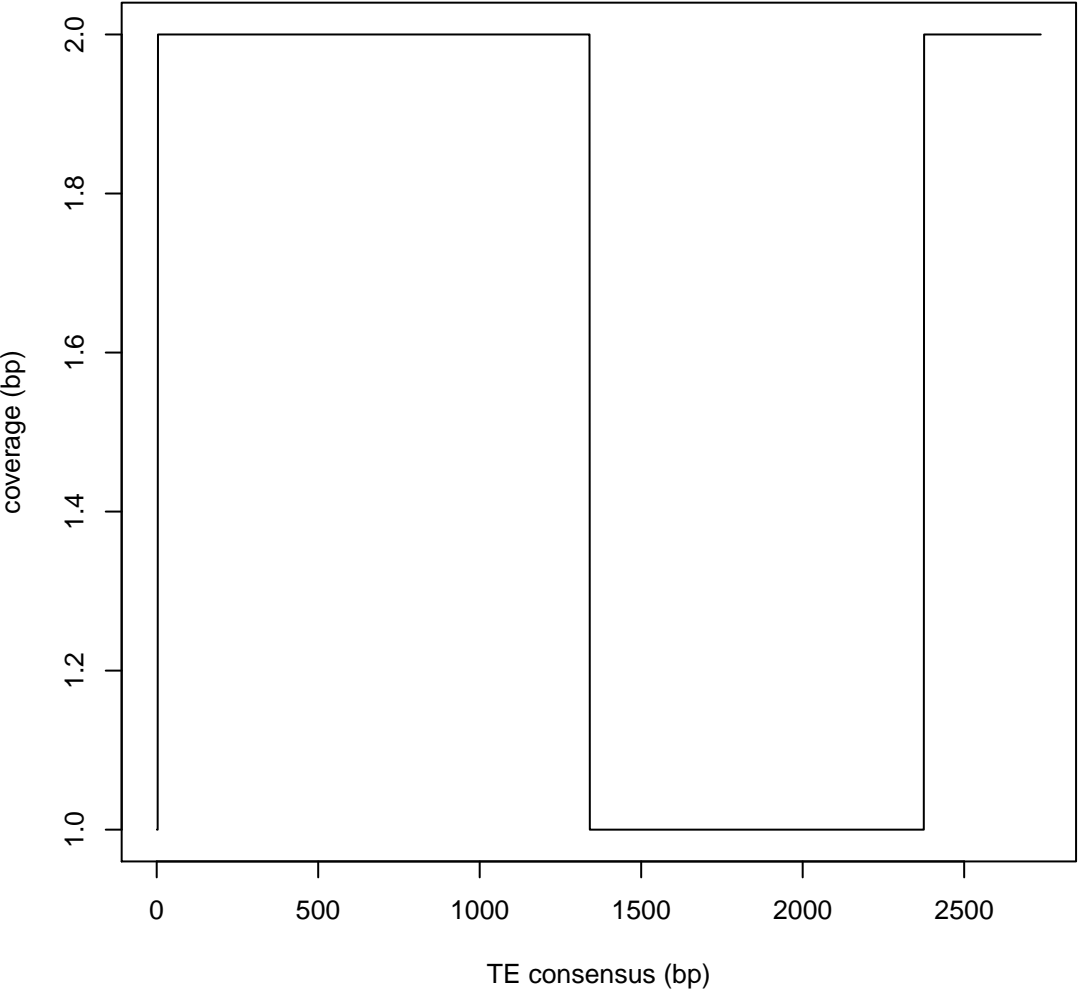


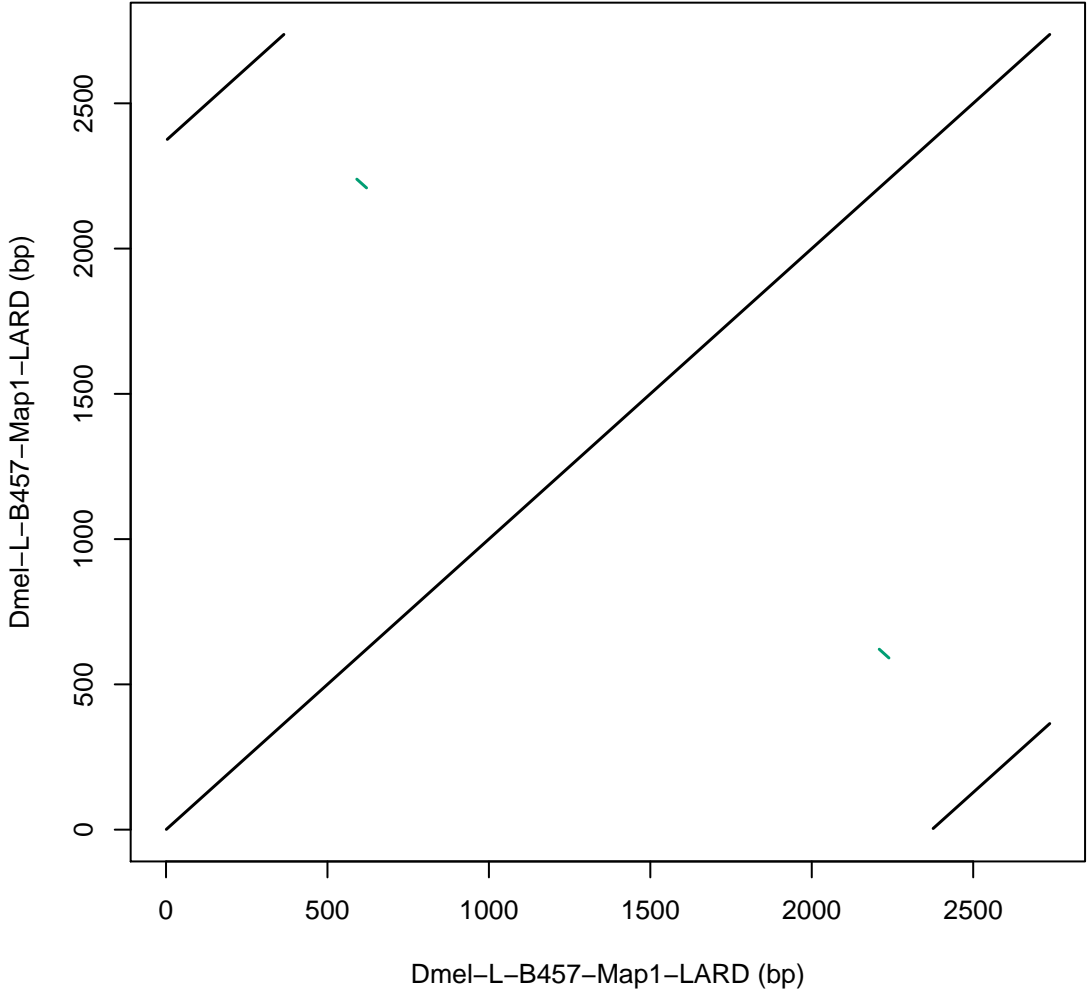
TE: Dmel-L-B457-Map1-LARD
consensus size: 2737bp; fragments: 3; full length: 1 (>=2463.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

