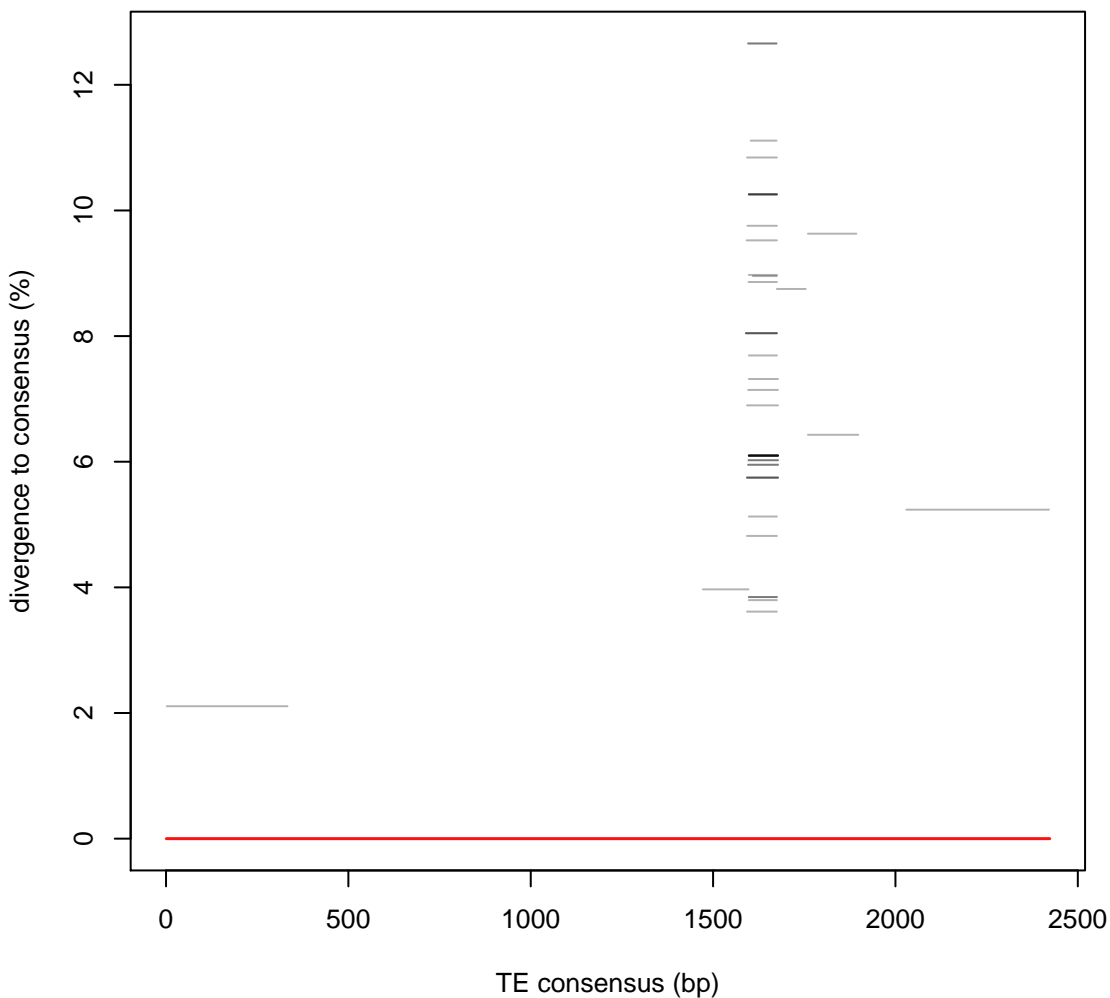
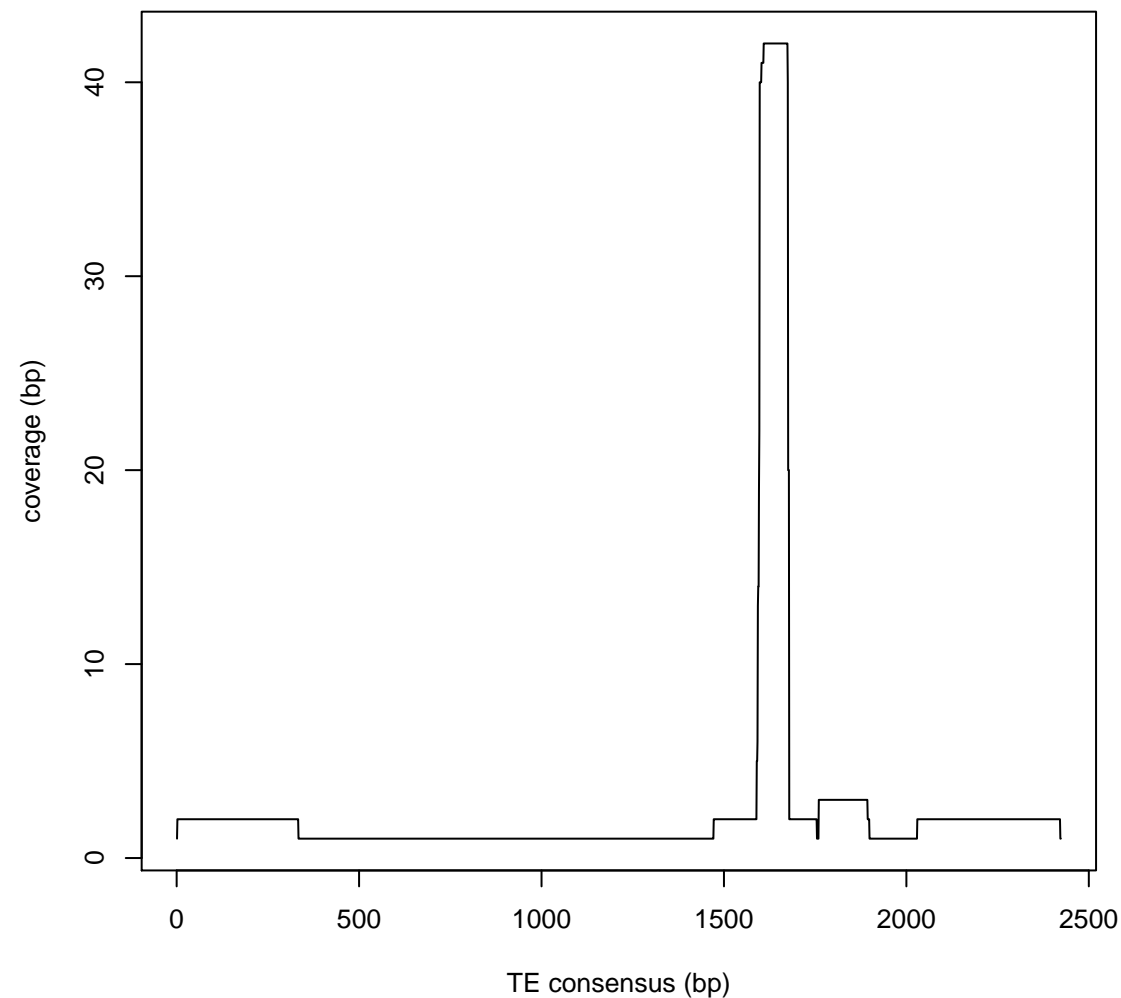


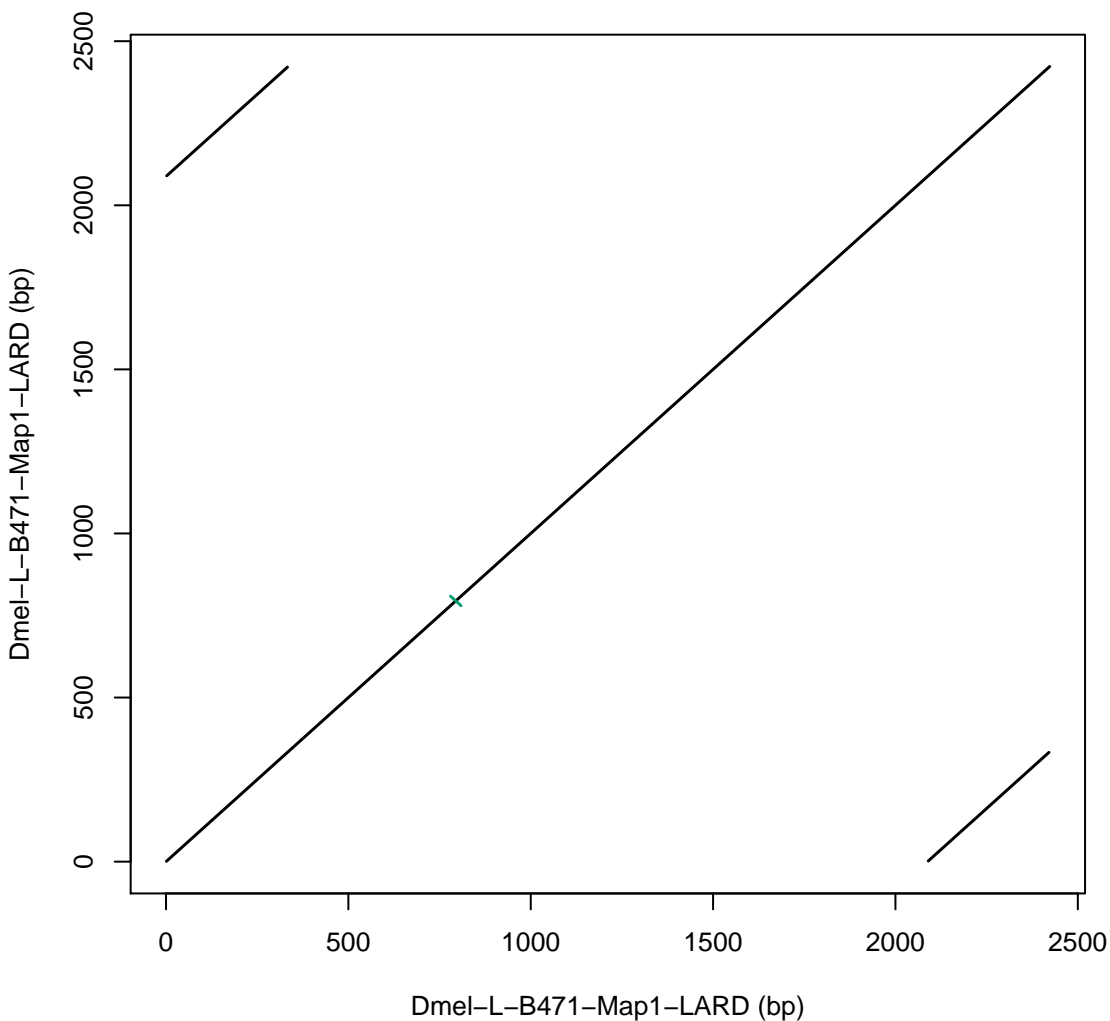
TE: Dmel-L-B471-Map1-LARD
consensus size: 2423bp; fragments: 48; full length: 1 (>=2180.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

