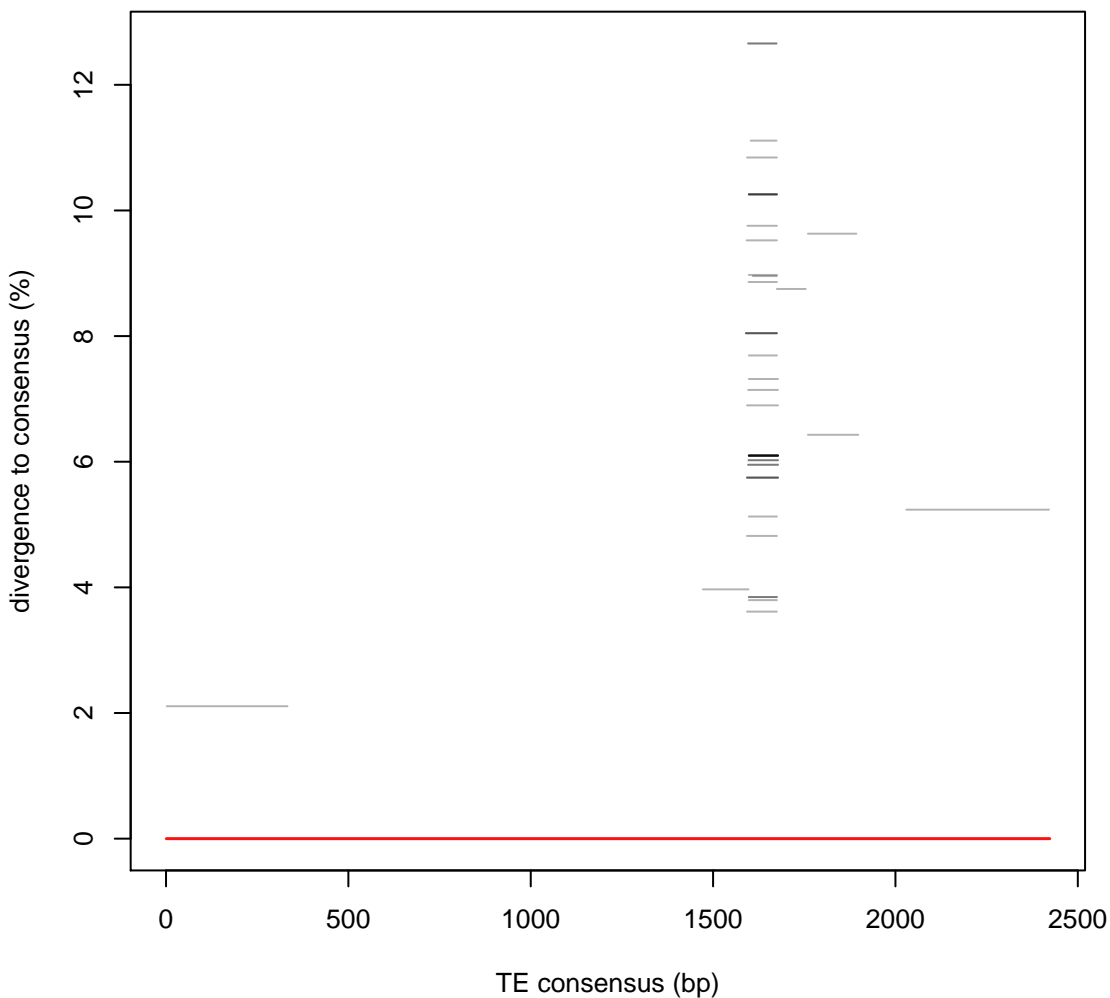
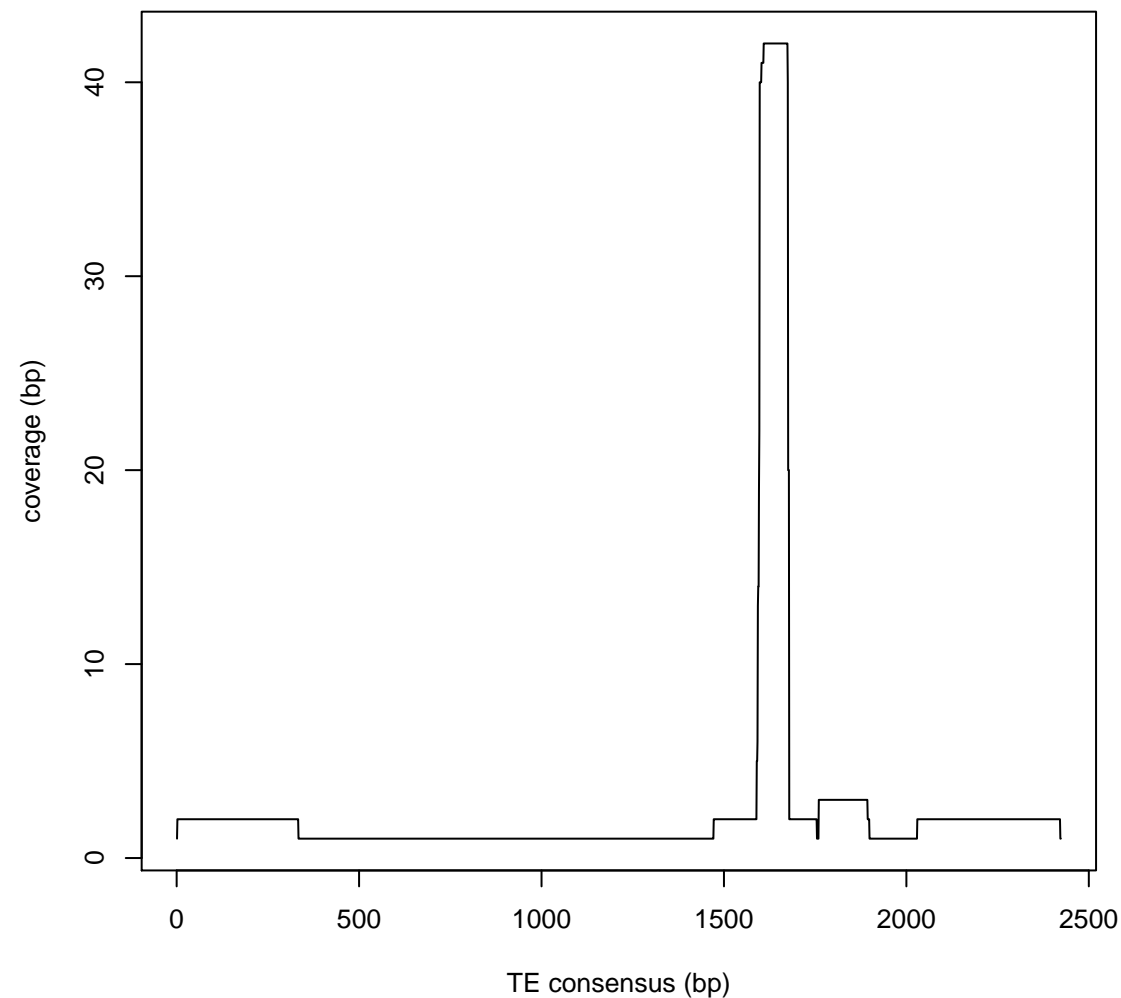


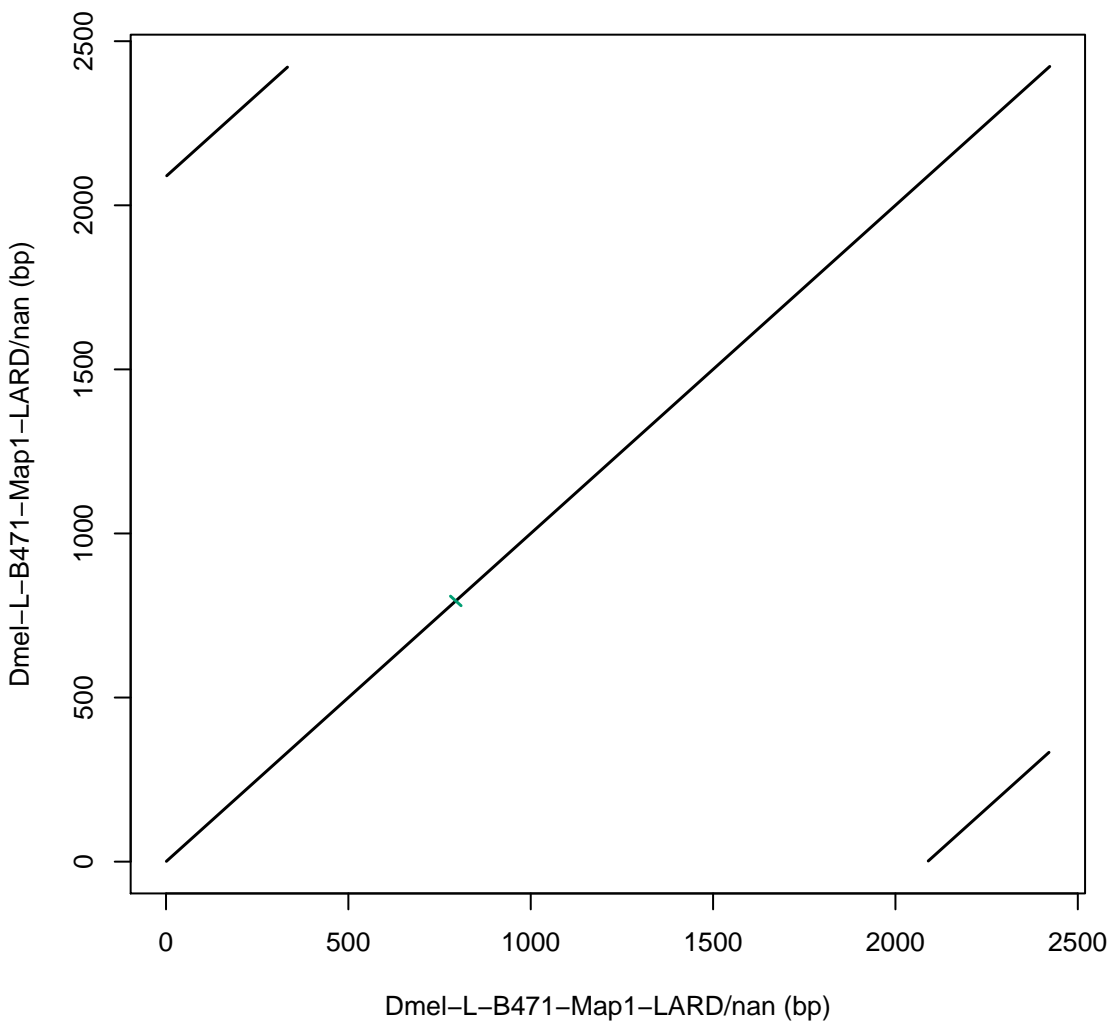
TE: Dmel-L-B471-Map1-LARD/nan
consensus size: 2423bp; fragments: 48; full length: 1 (≥ 2180.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

