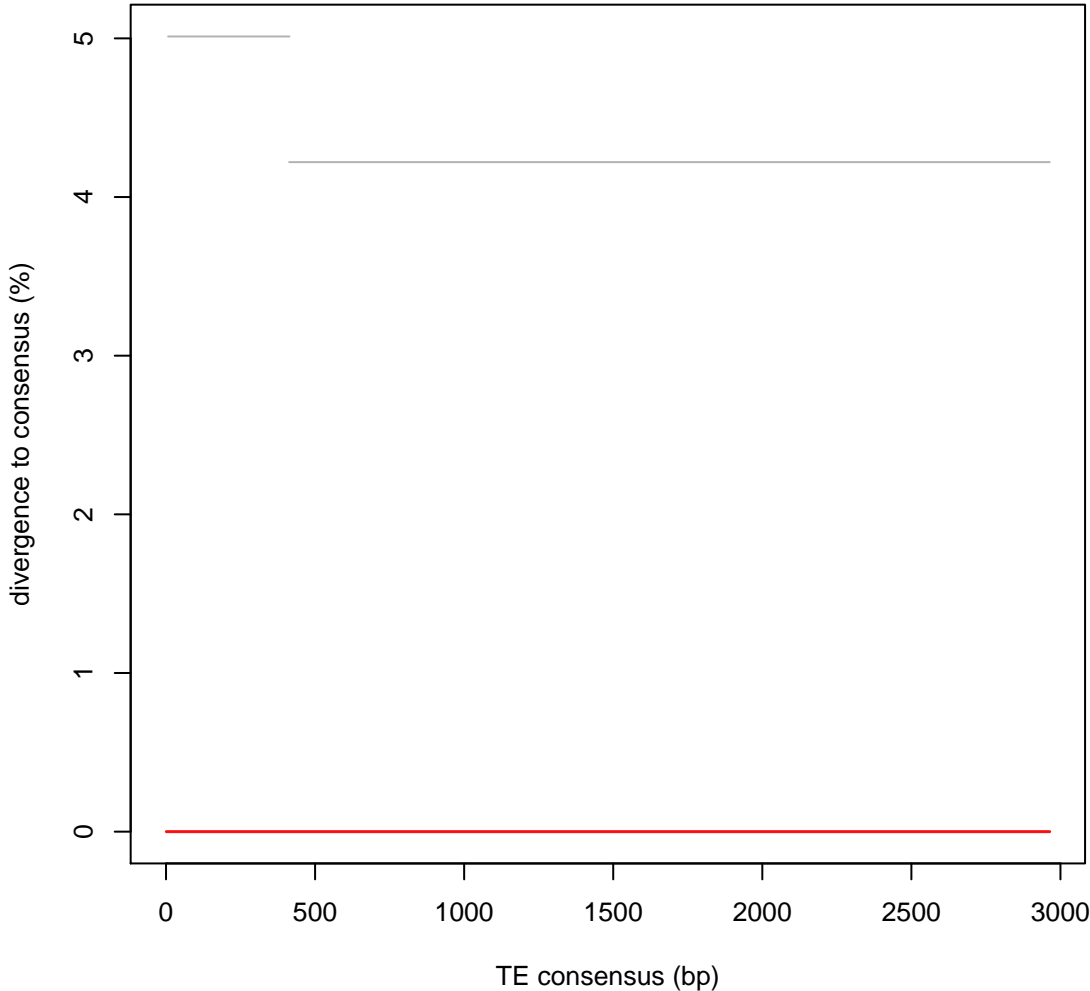
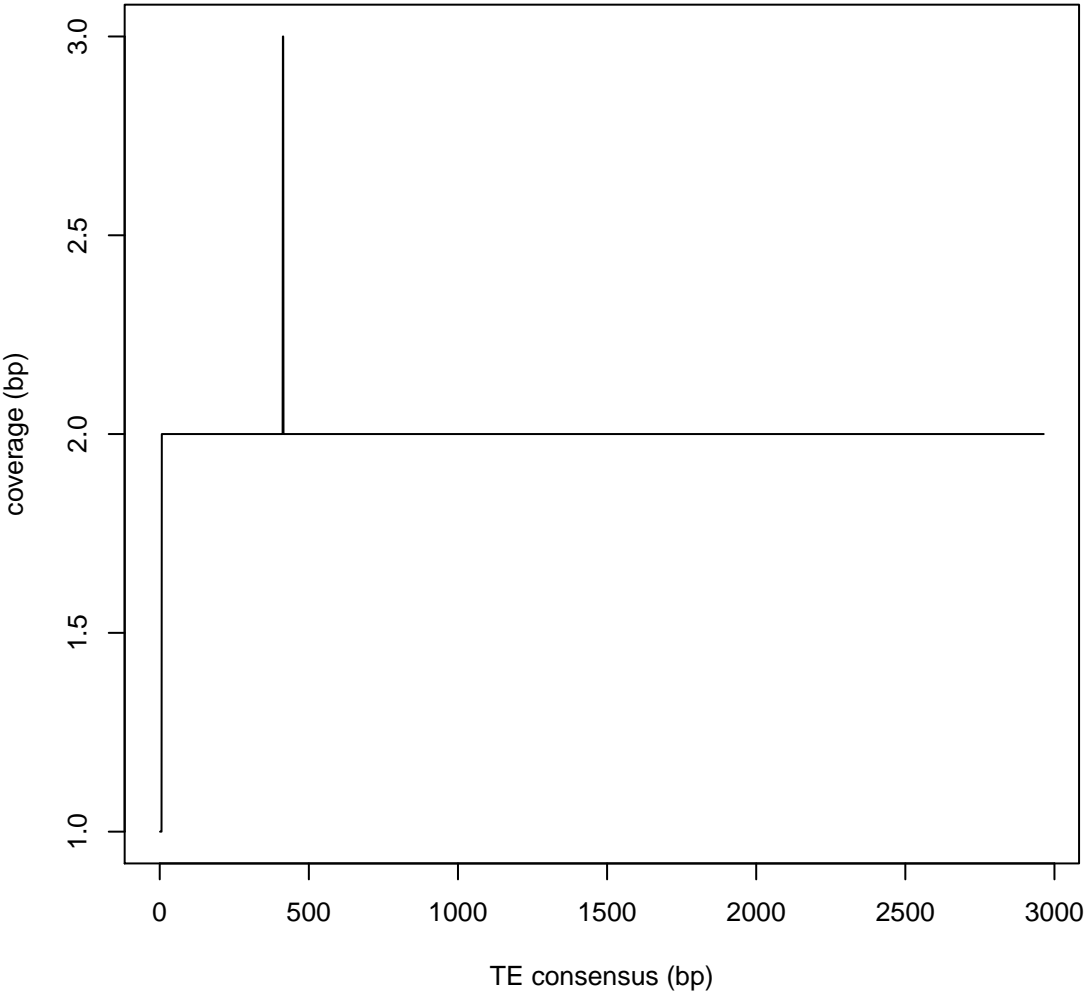


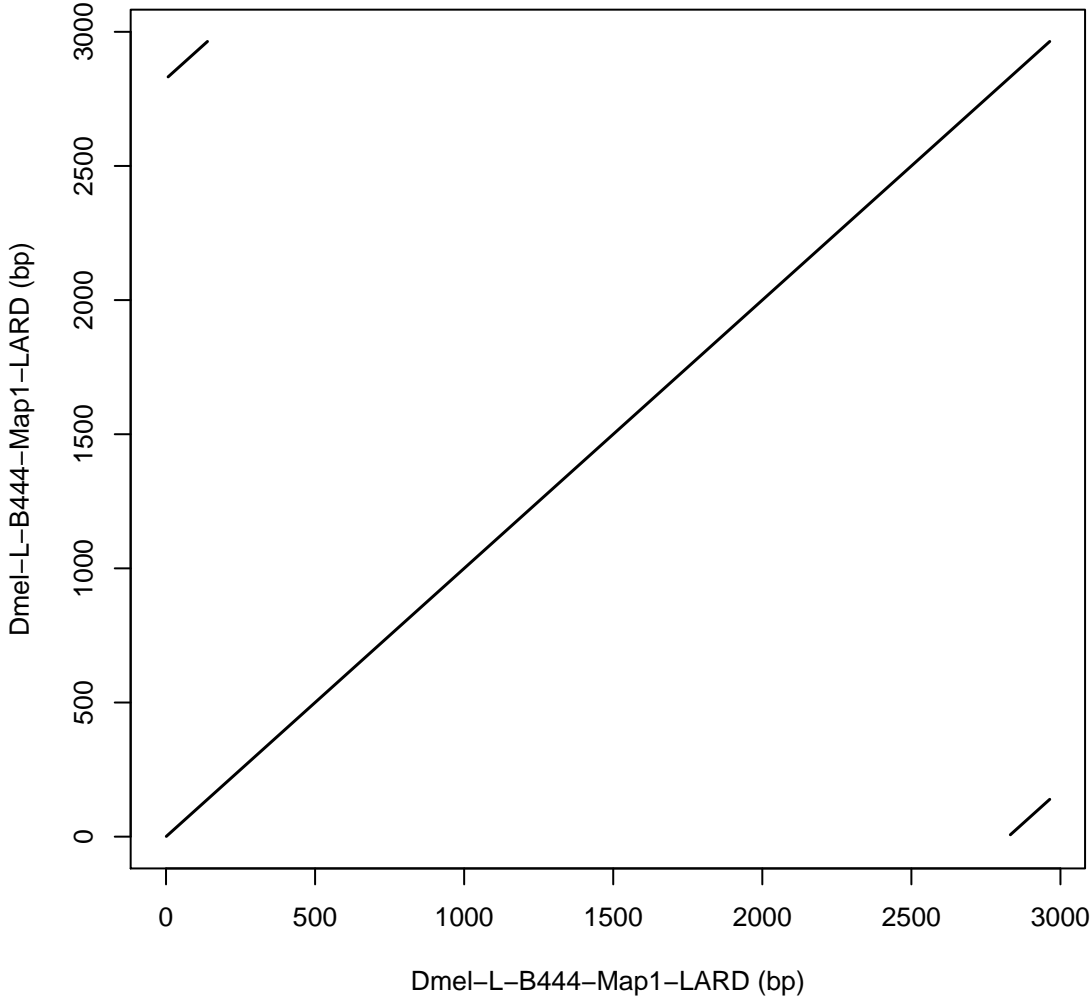
TE: Dmel-L-B444-Map1-LARD
consensus size: 2964bp; fragments: 3; full length: 1 (>=2667.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

