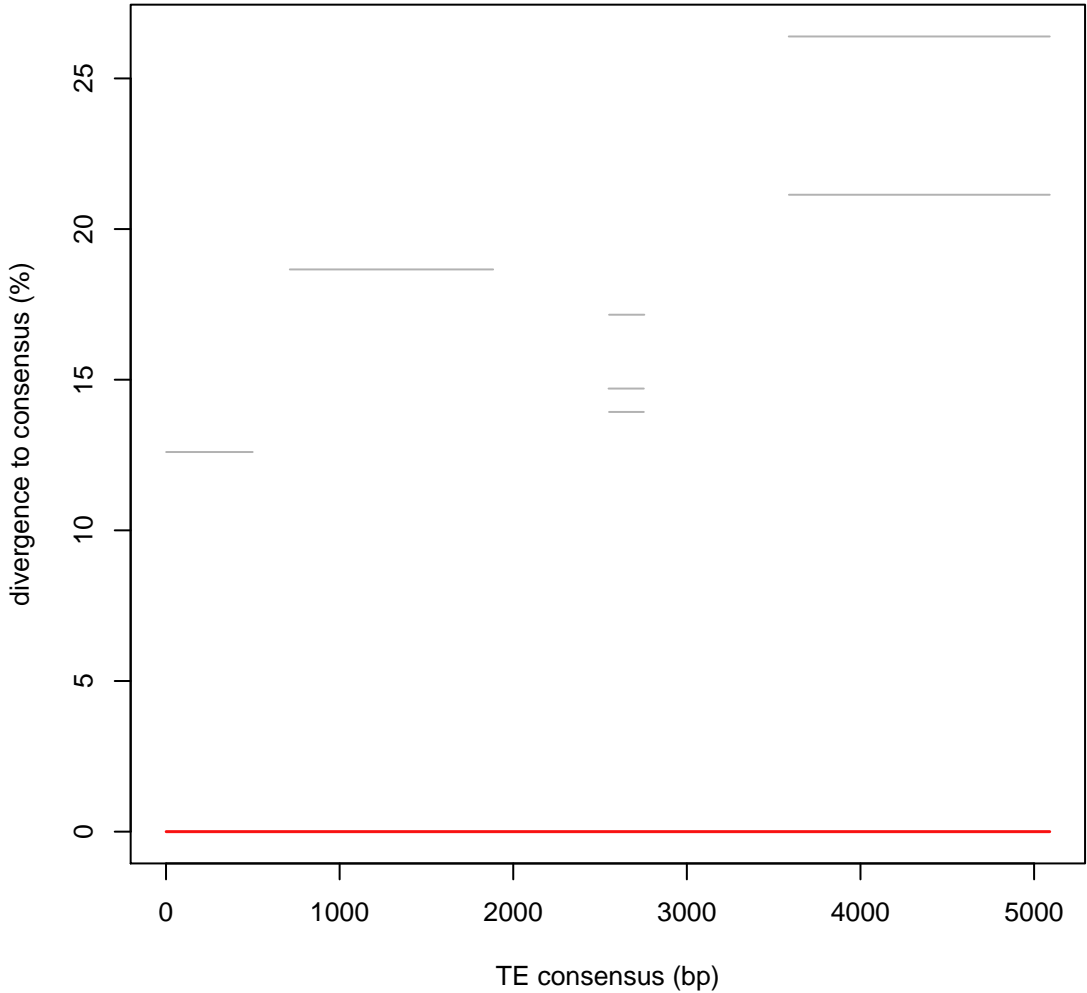
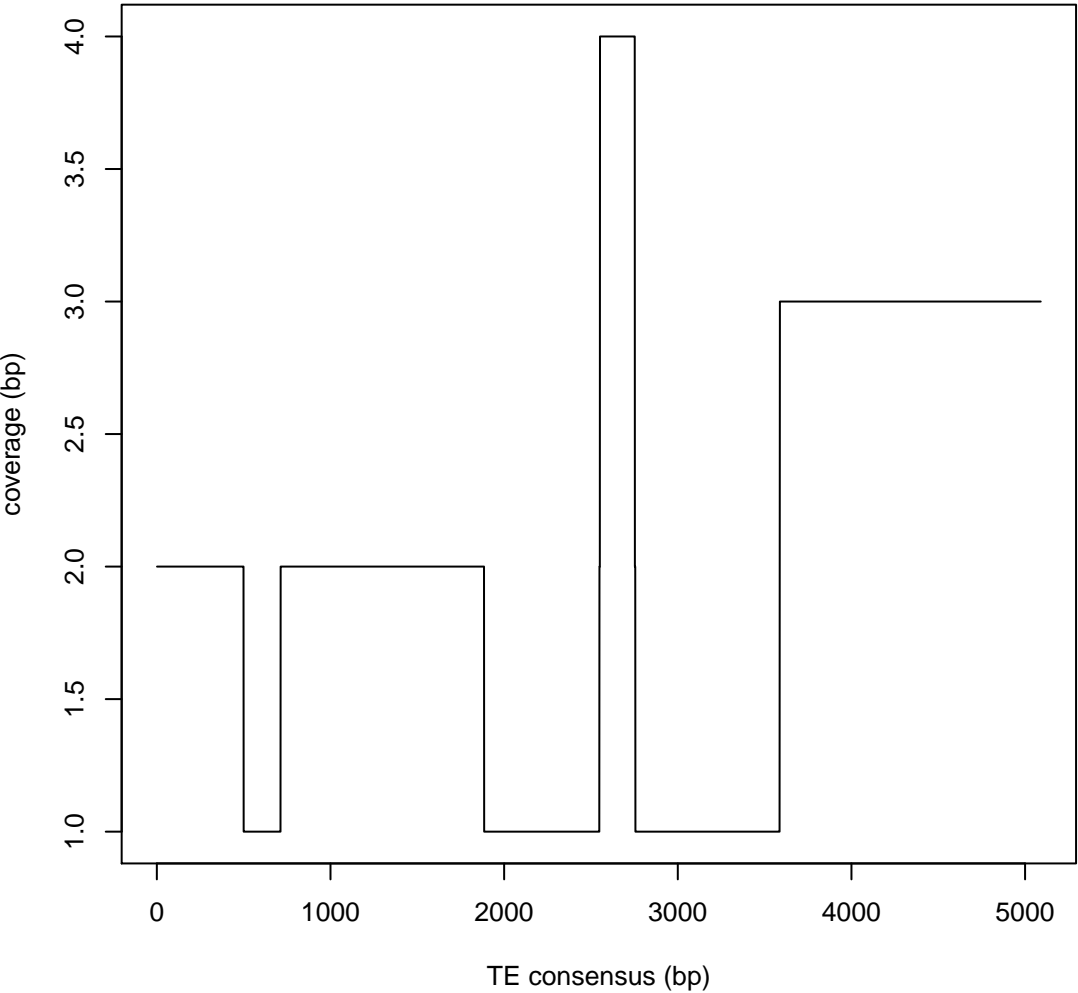


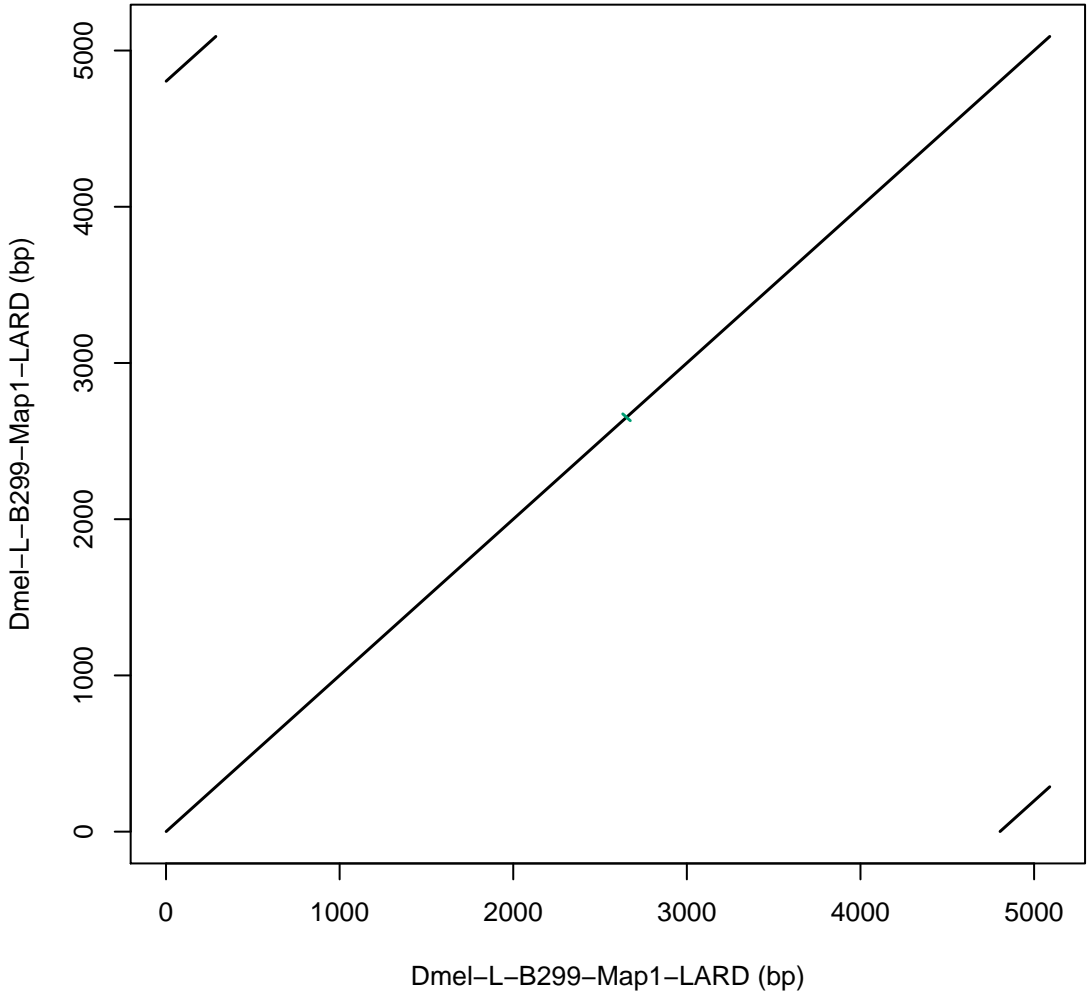
TE: Dmel-L-B299-Map1-LARD
consensus size: 5090bp; fragments: 8; full length: 1 (>=4581bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

