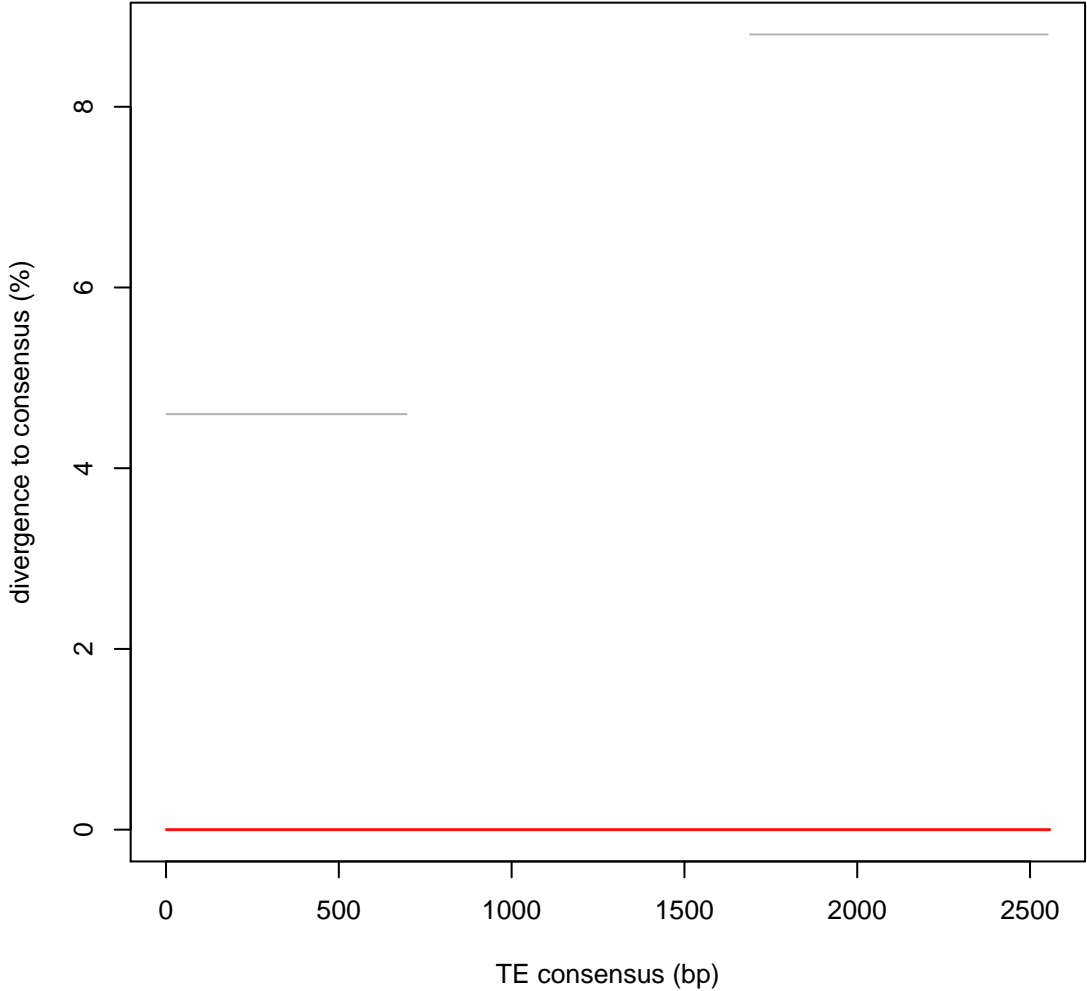
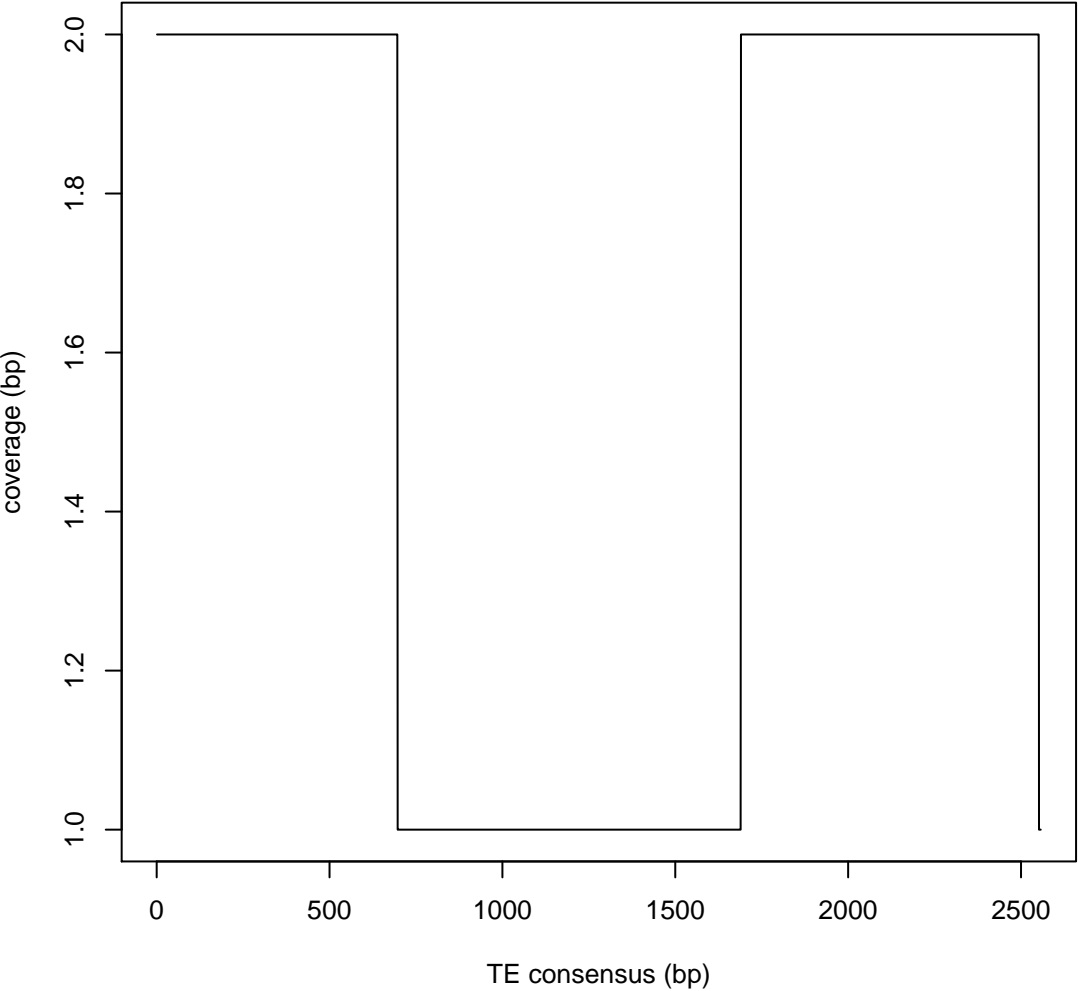


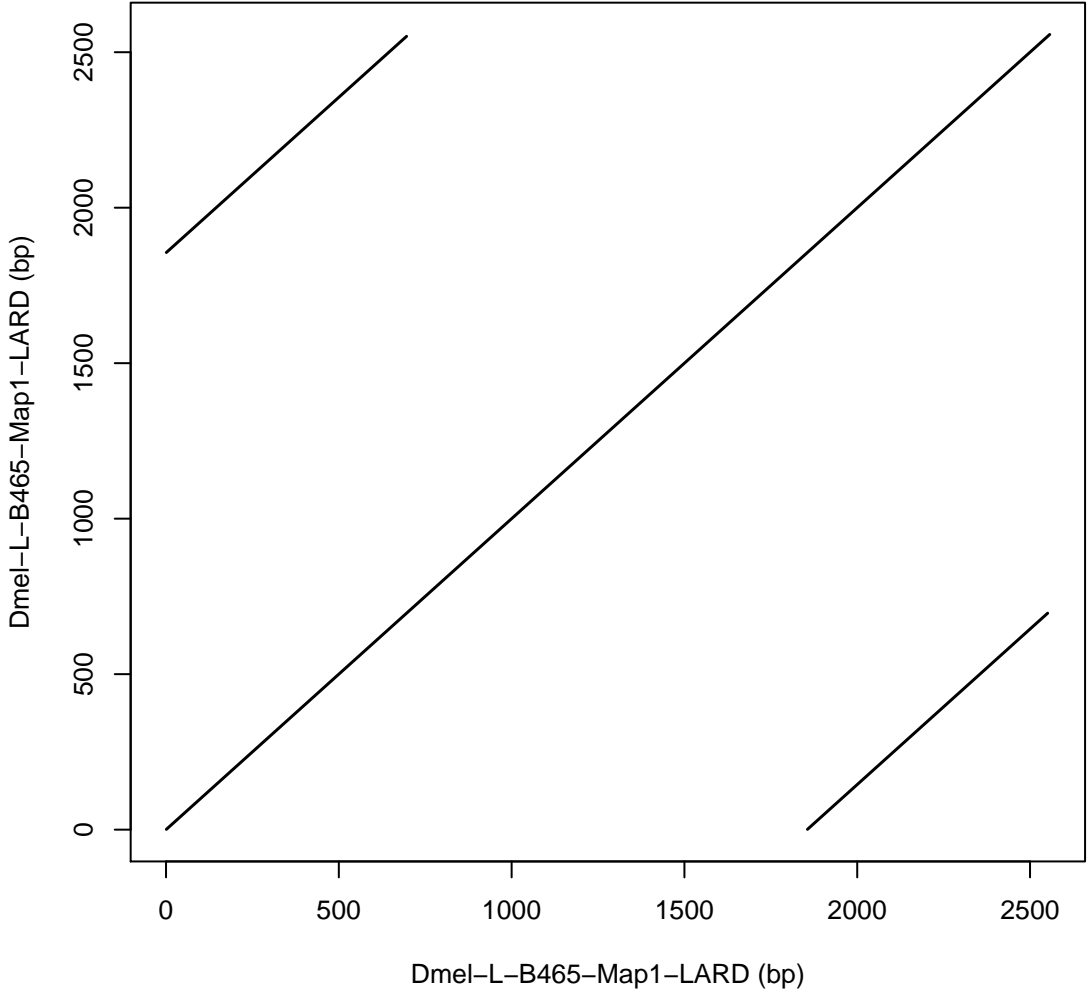
TE: Dmel-L-B465-Map1-LARD
consensus size: 2557bp; fragments: 3; full length: 1 (>=2301.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

