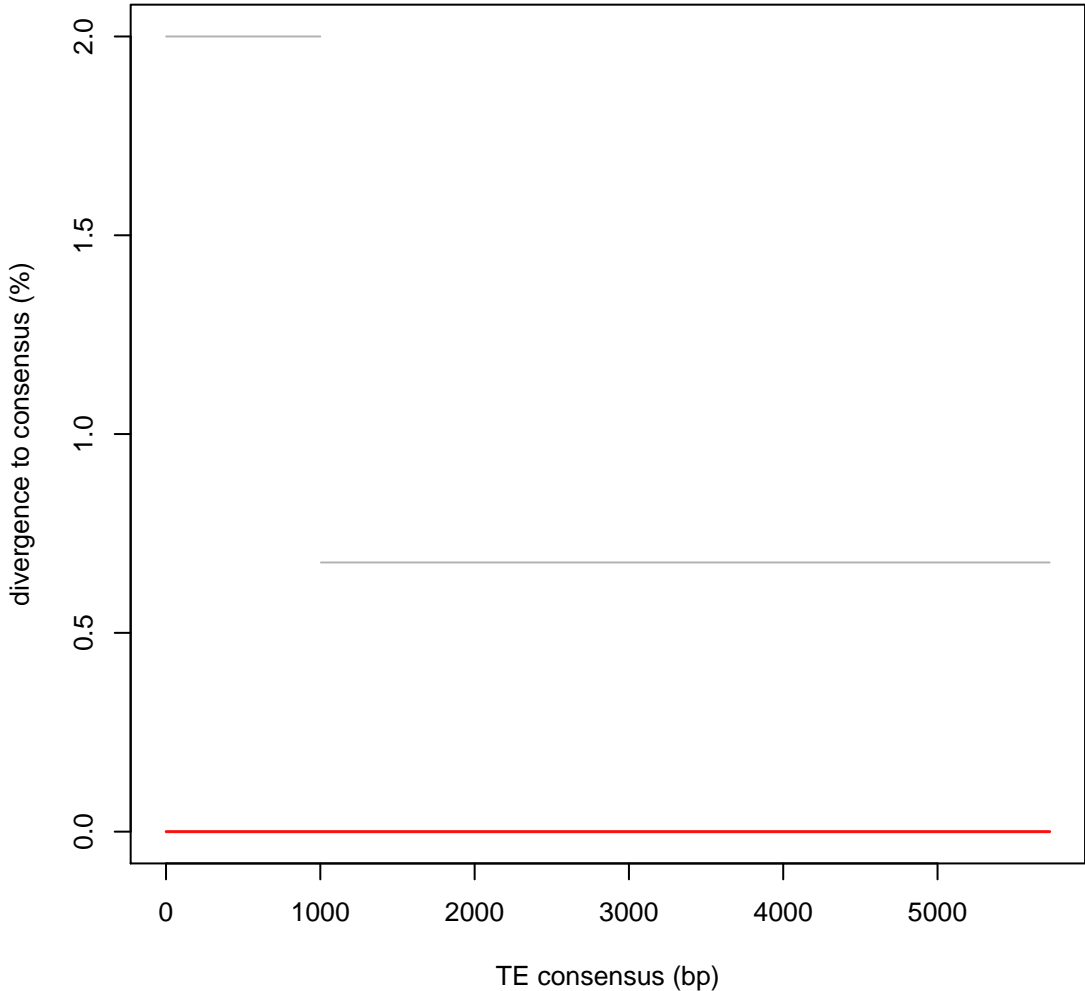
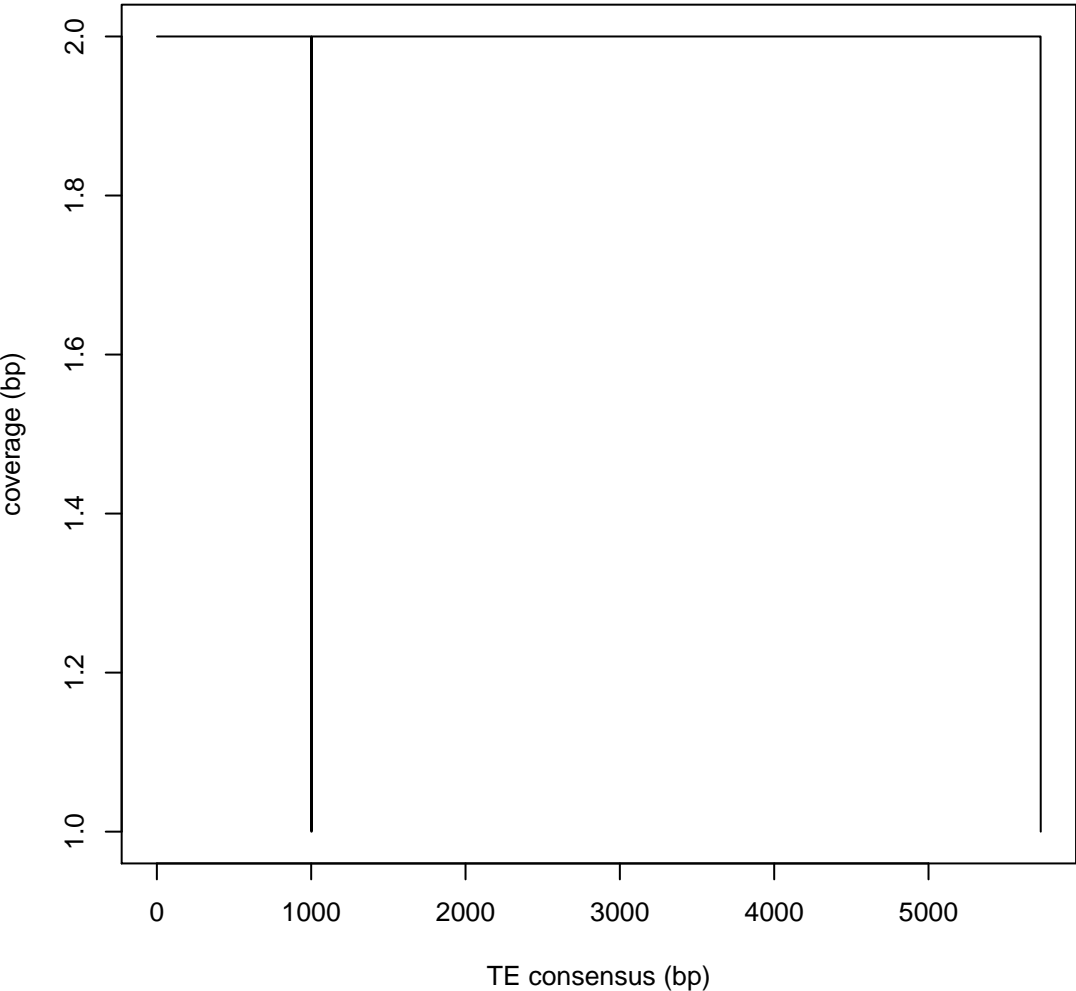


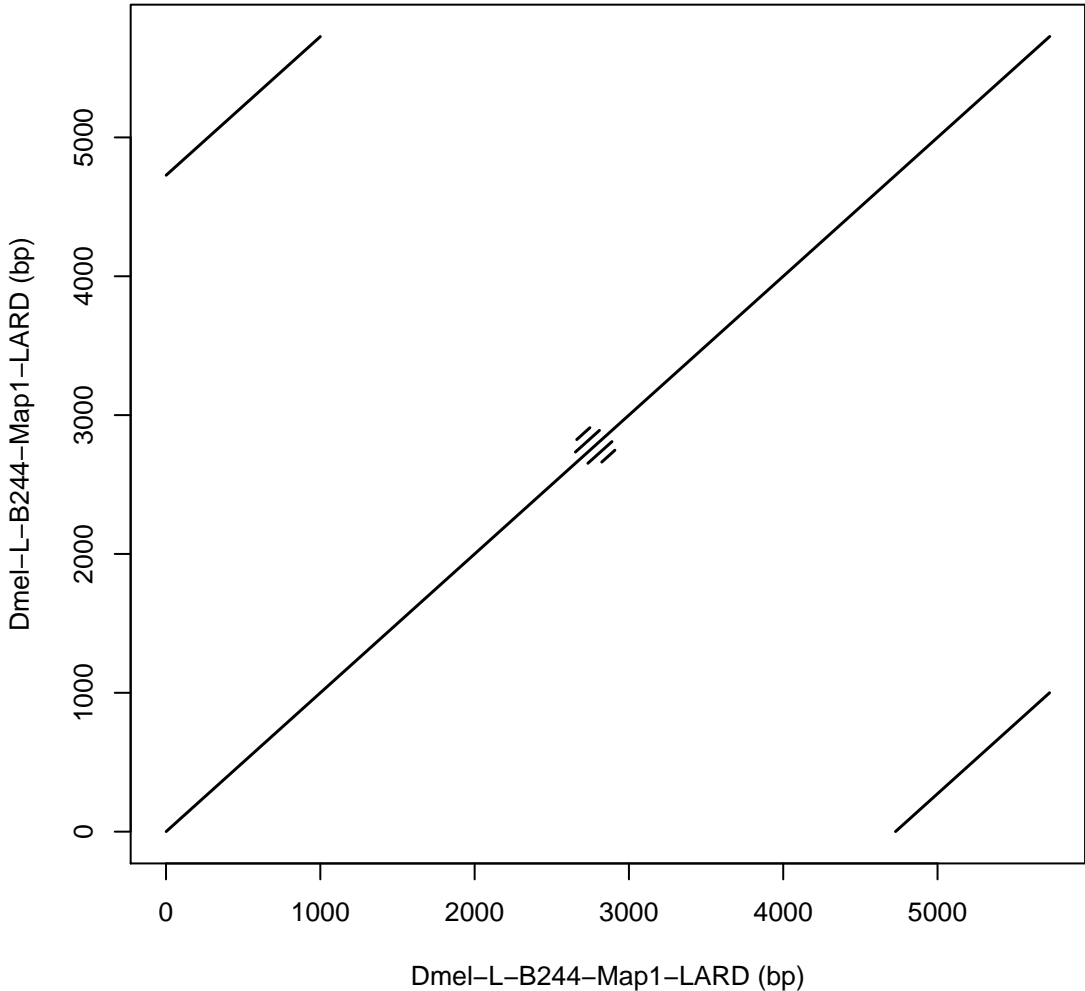
TE: Dmel-L-B244-Map1-LARD
consensus size: 5727bp; fragments: 3; full length: 1 (>=5154.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

