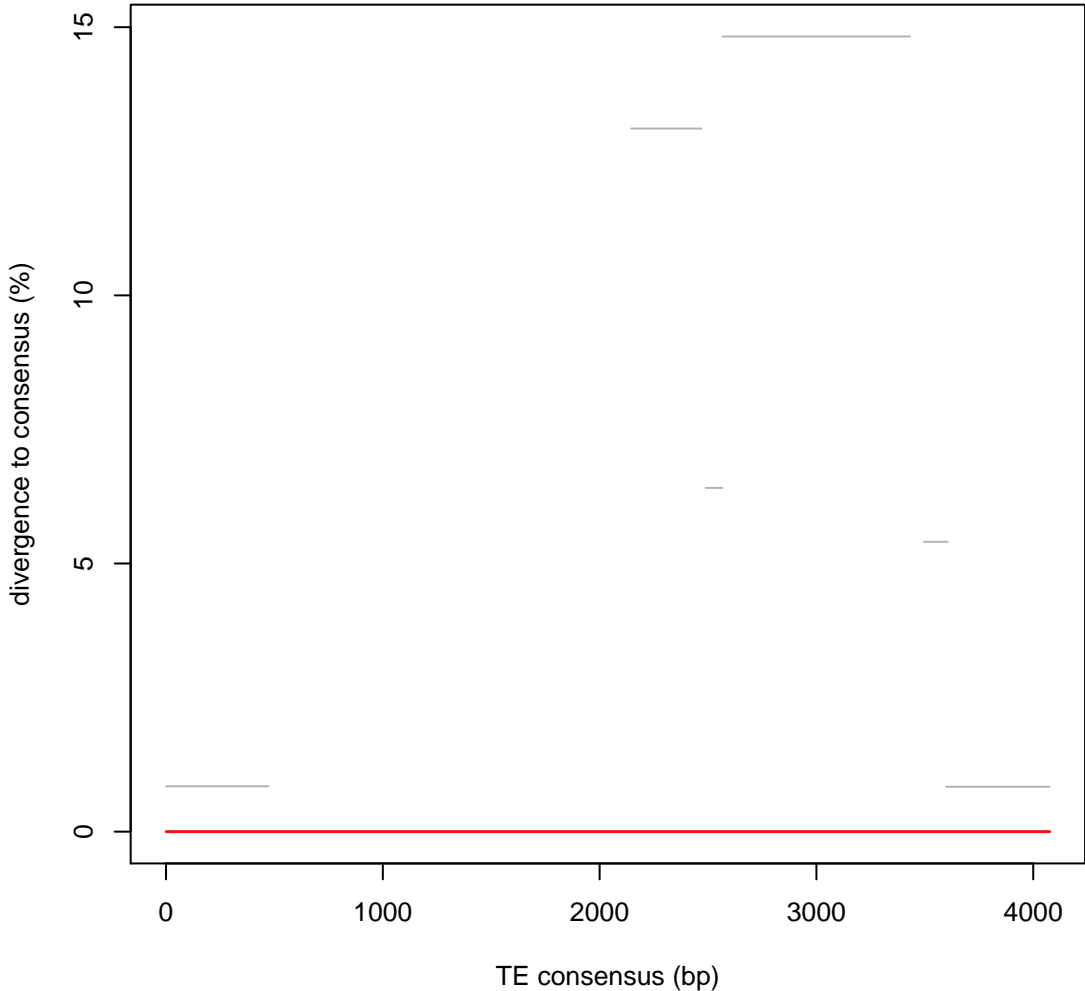
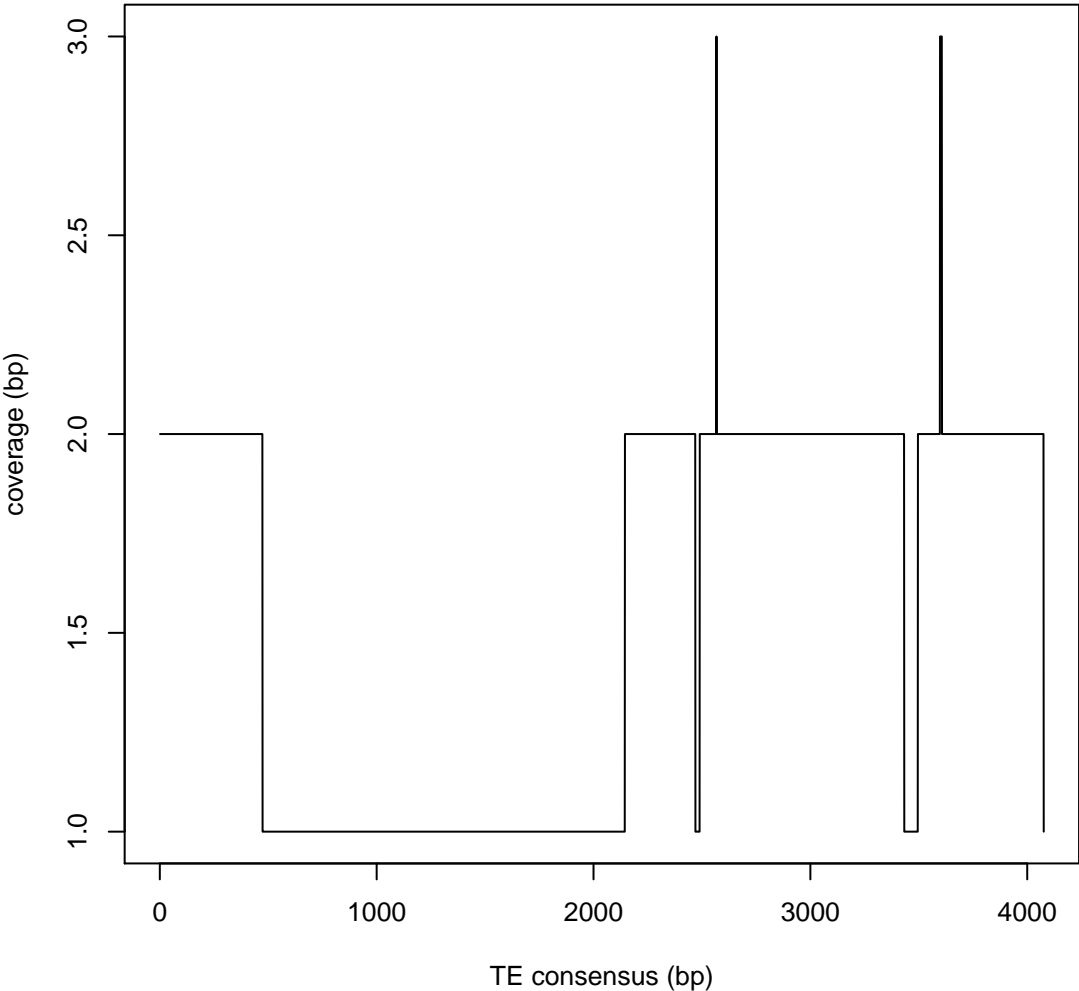


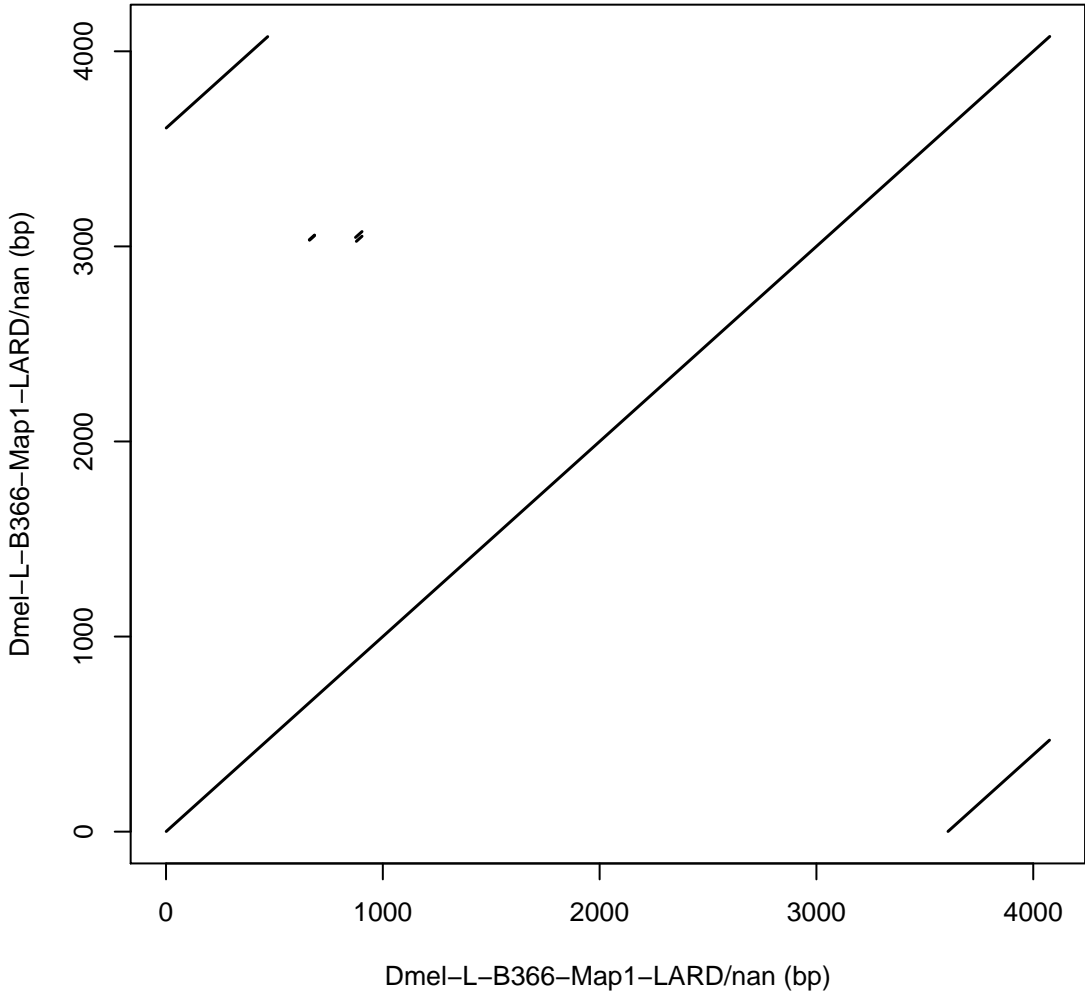
TE: Dmel-L-B366-Map1-LARD/nan
consensus size: 4076bp; fragments: 7; full length: 1 (>=3668.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

