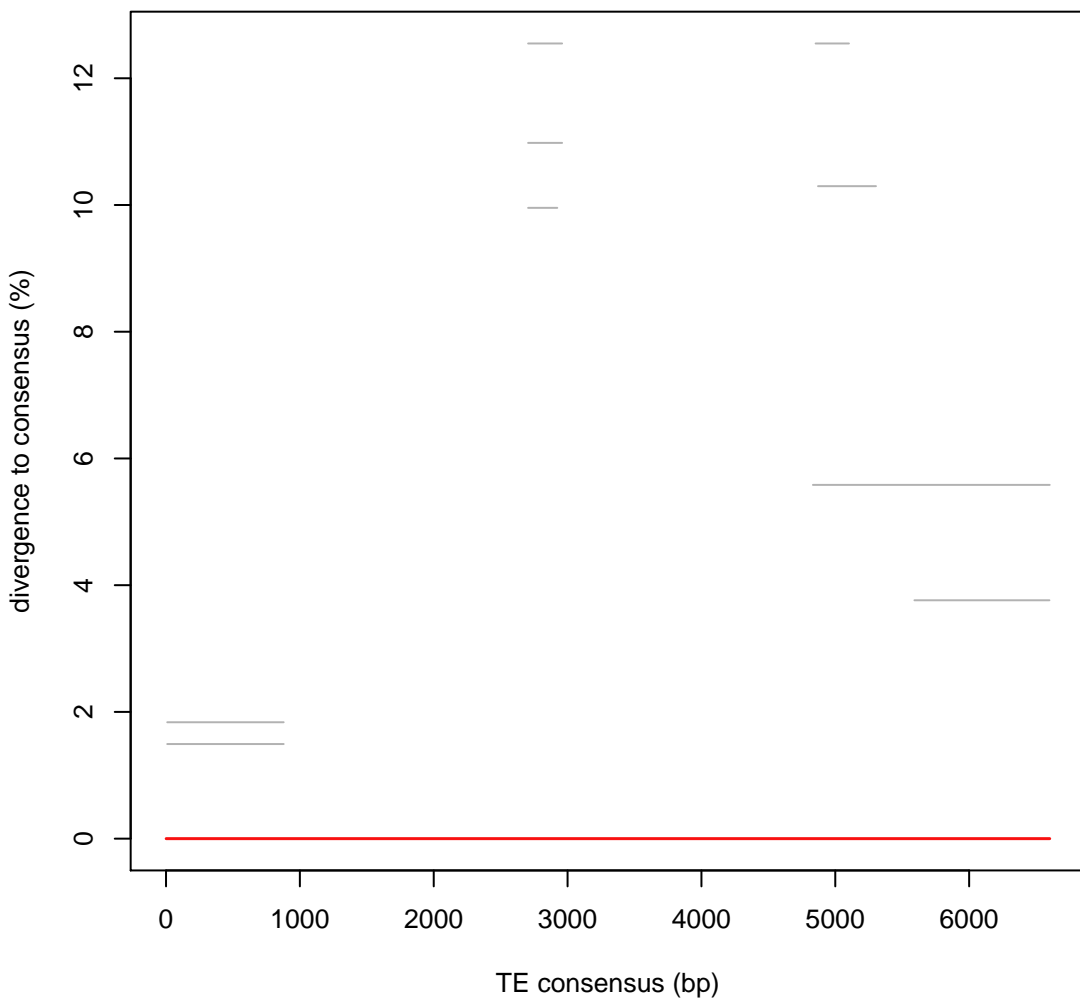
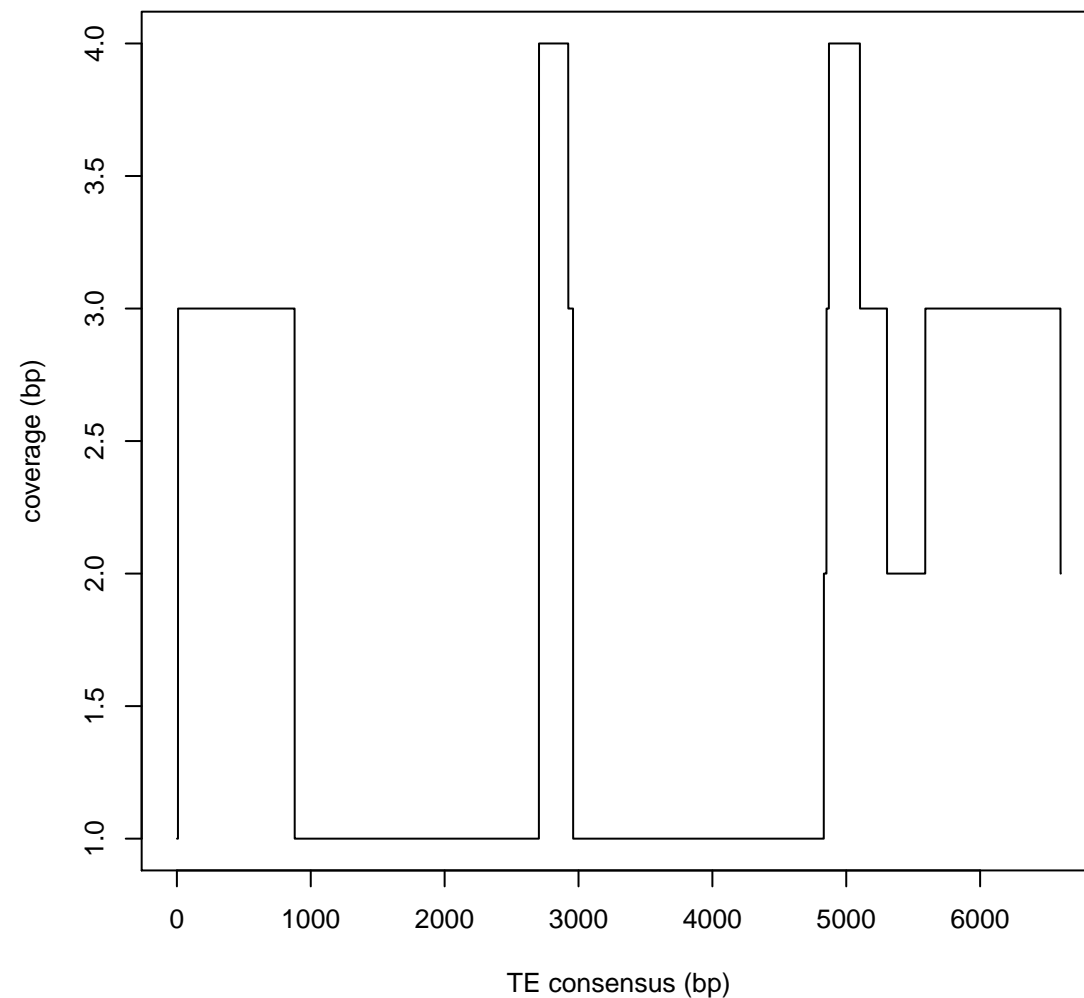


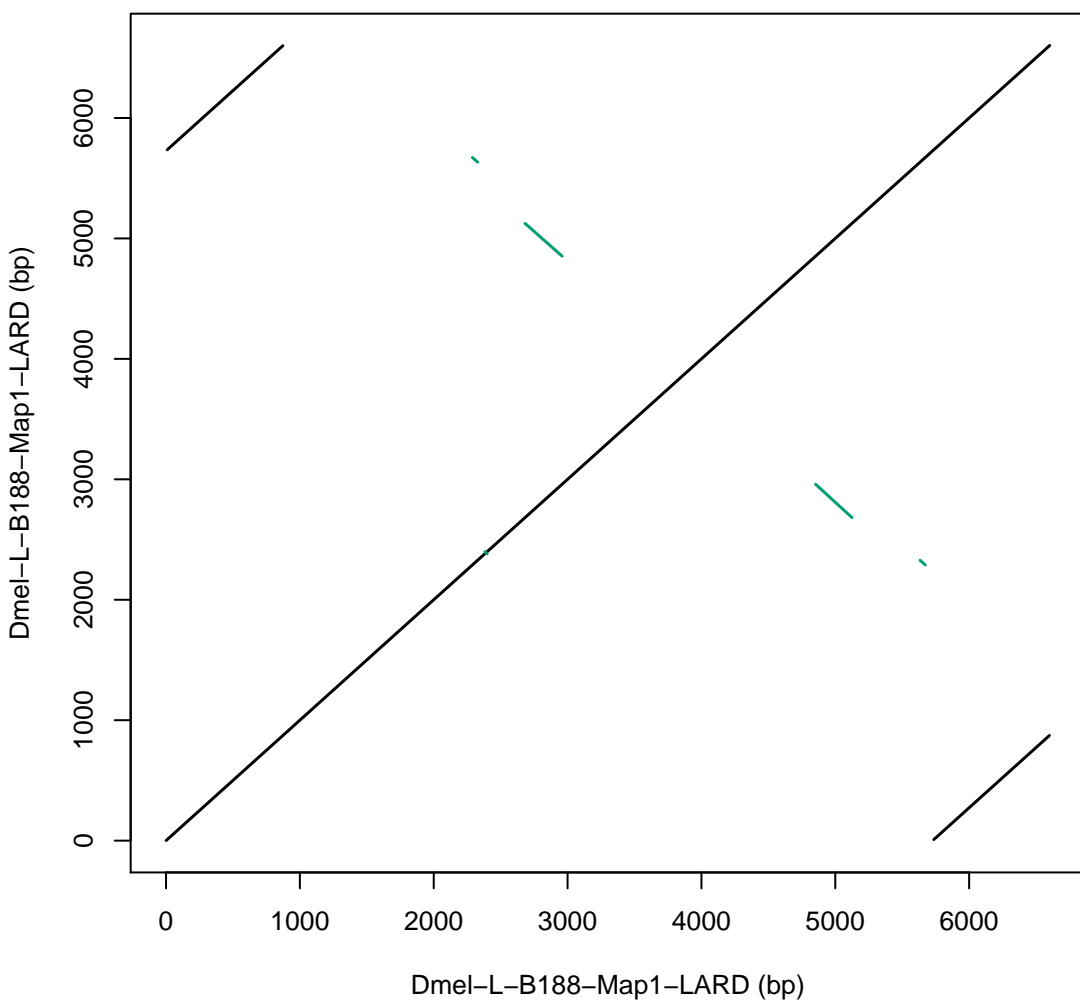
TE: Dmel-L-B188-Map1-LARD
consensus size: 6602bp; fragments: 10; full length: 1 (>=5941.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

