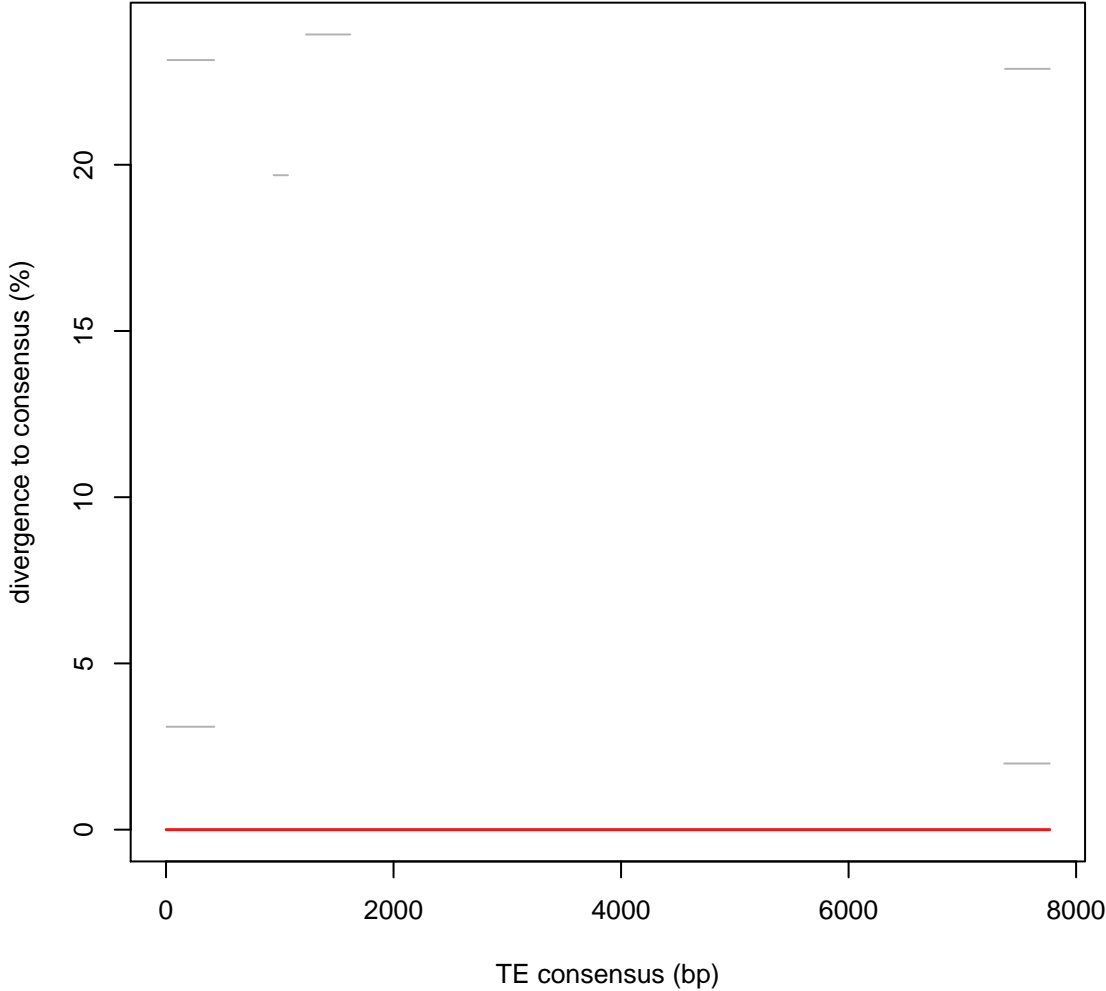
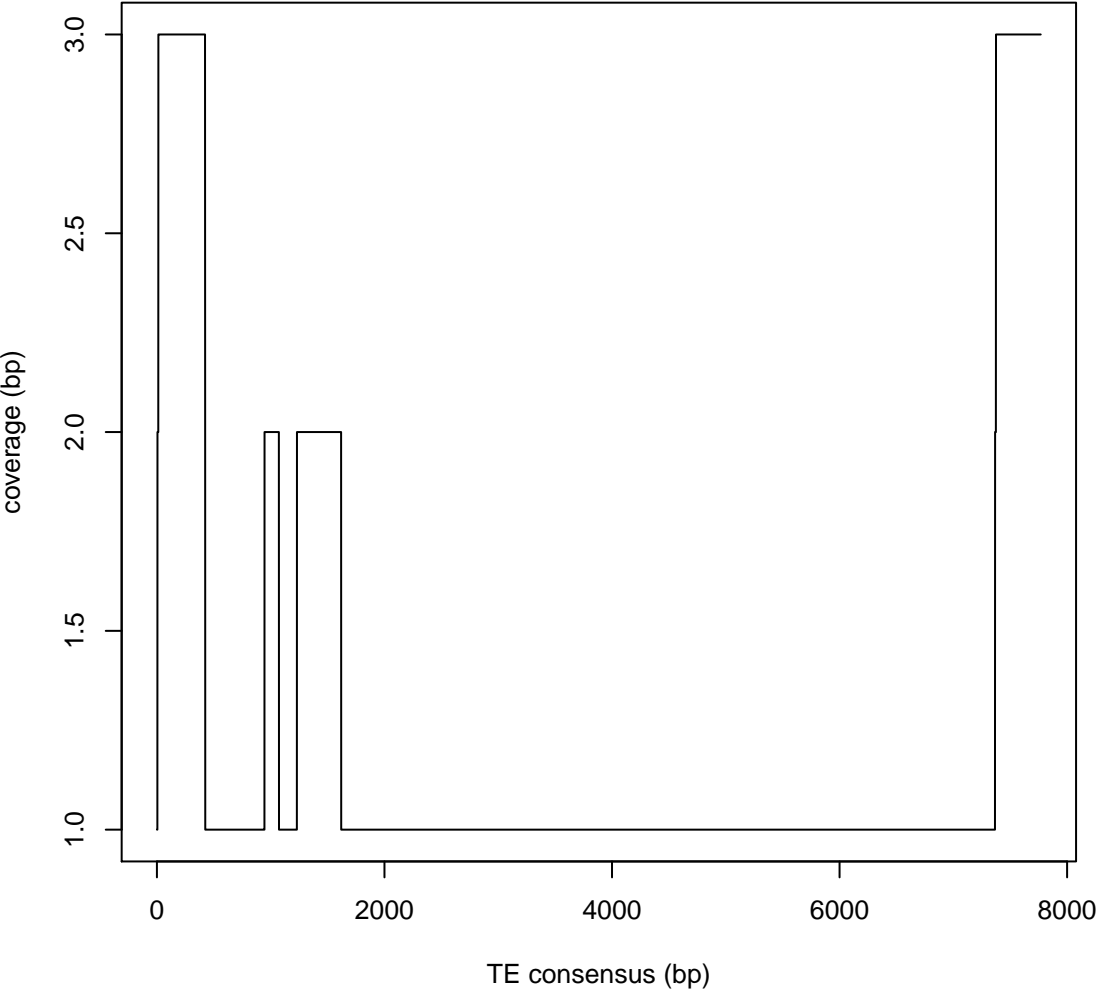


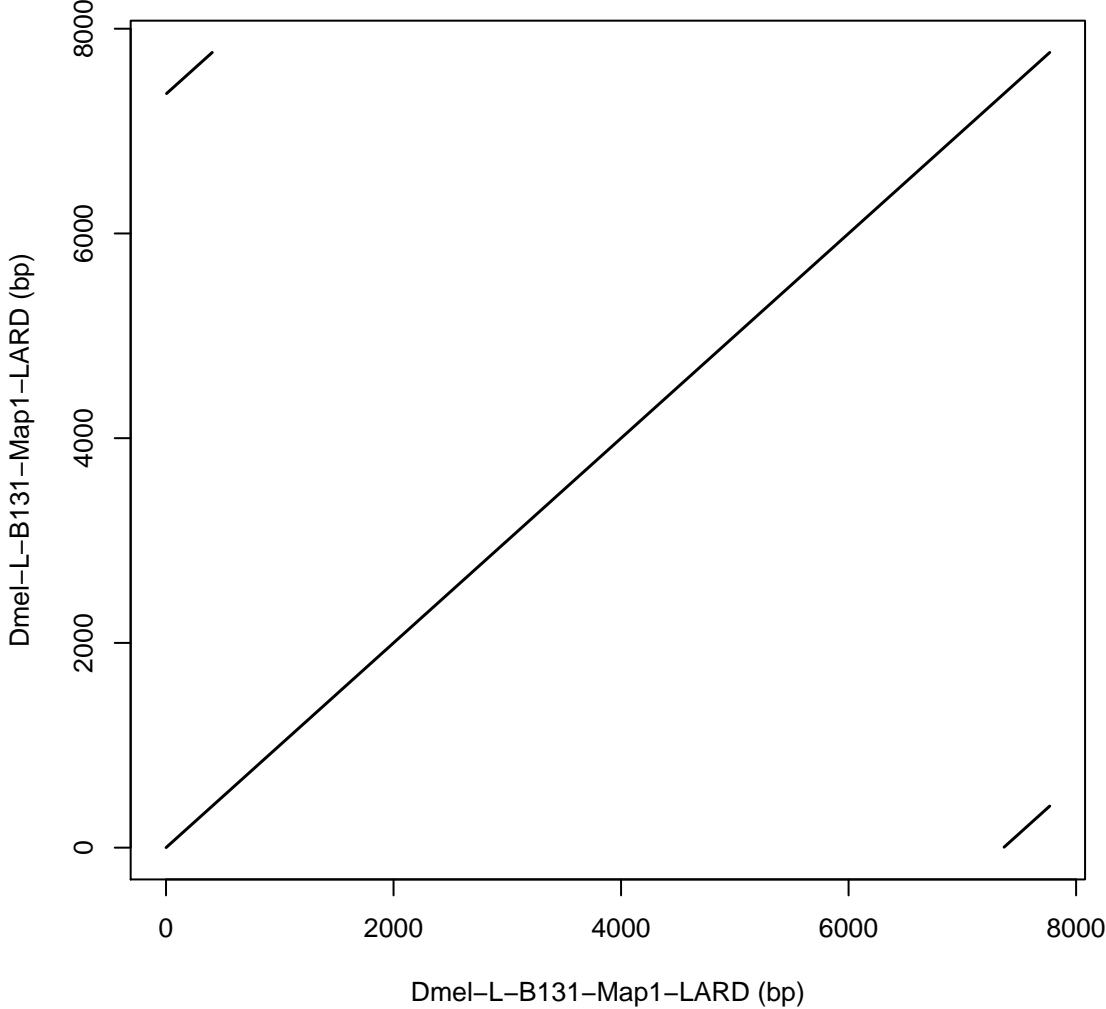
TE: Dmel-L-B131-Map1-LARD
consensus size: 7768bp; fragments: 7; full length: 1 (>=6991.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

