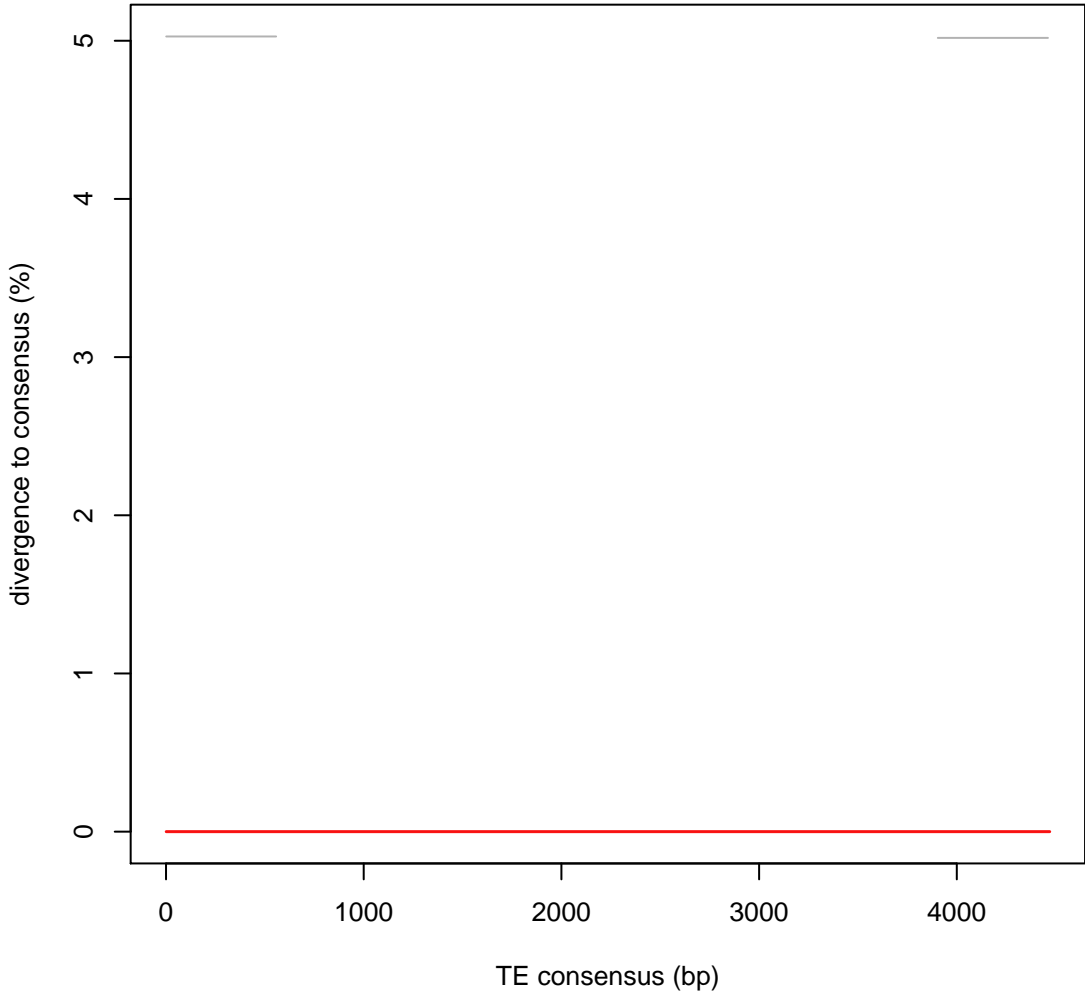
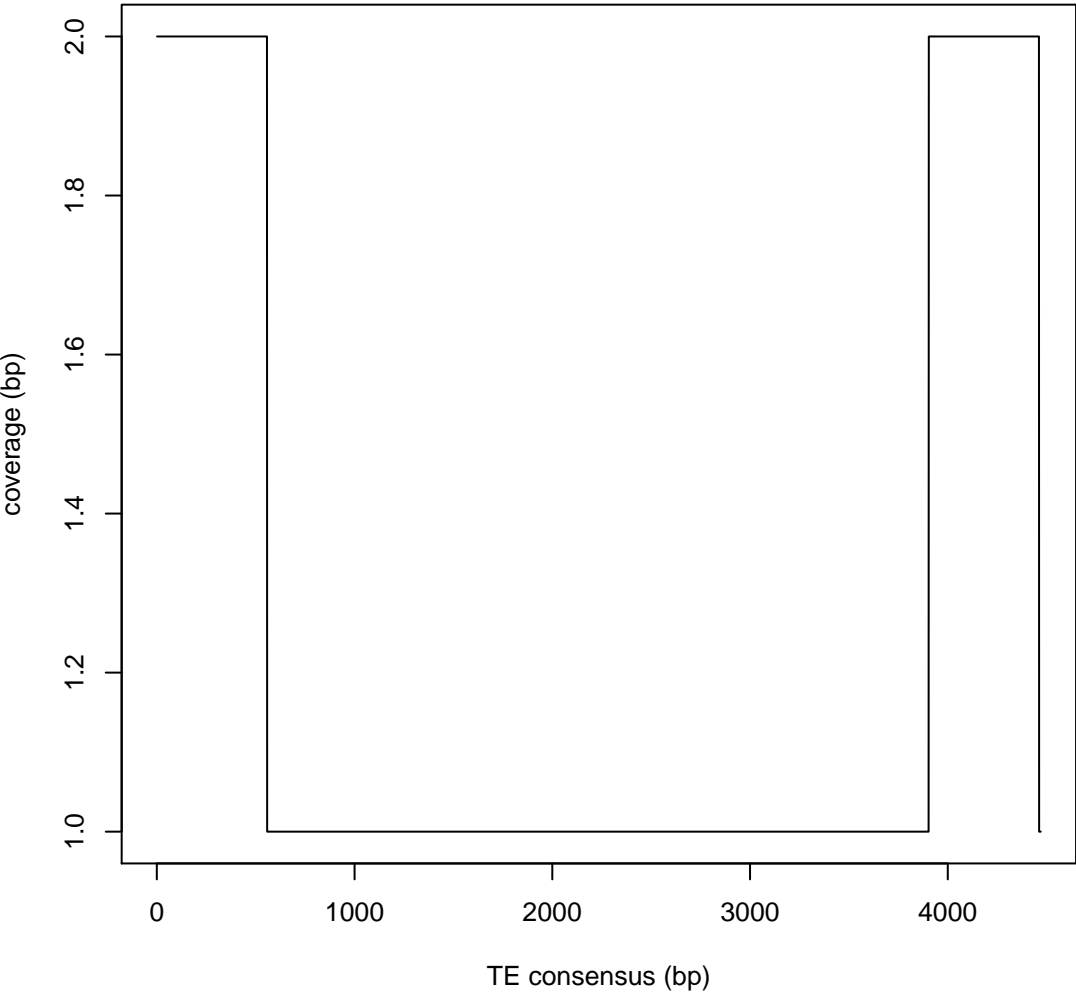


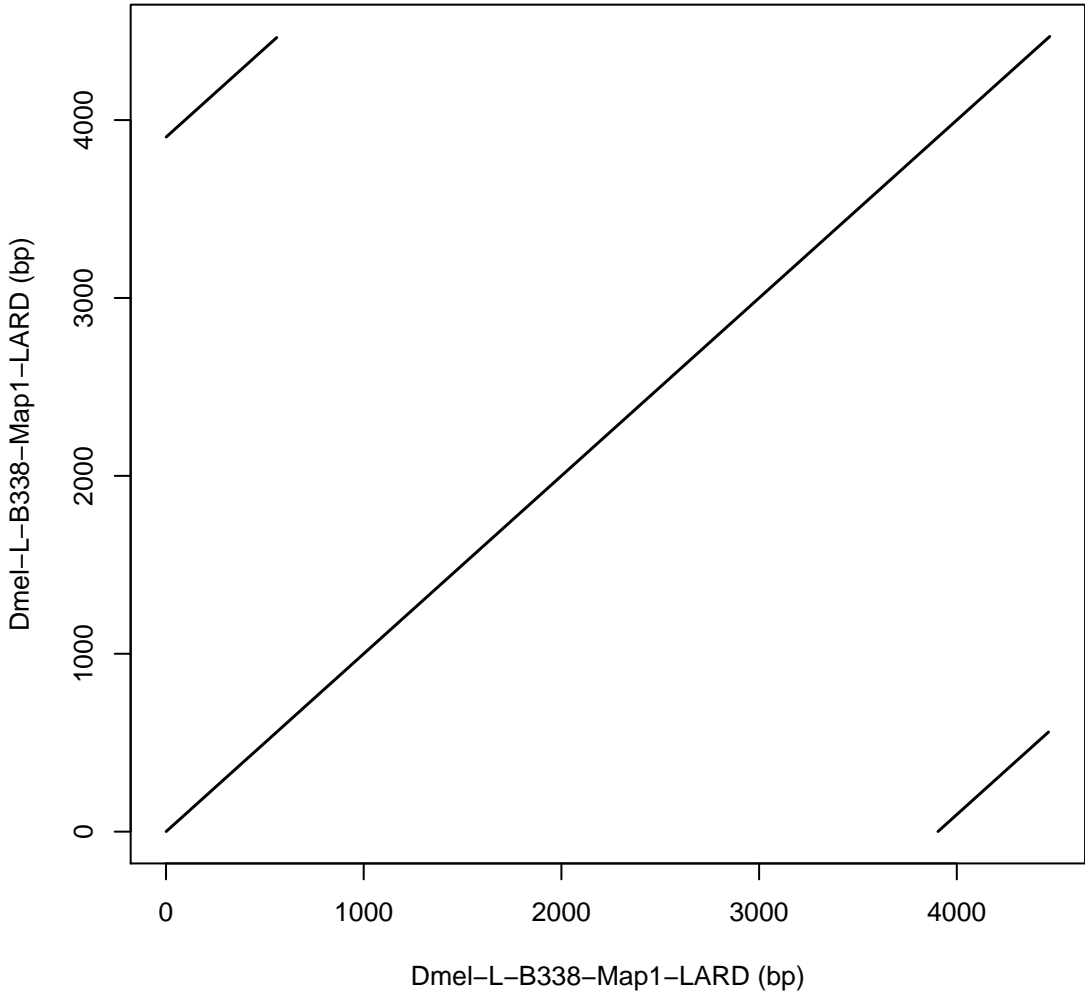
TE: Dmel-L-B338-Map1-LARD
consensus size: 4470bp; fragments: 3; full length: 1 (>=4023bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

