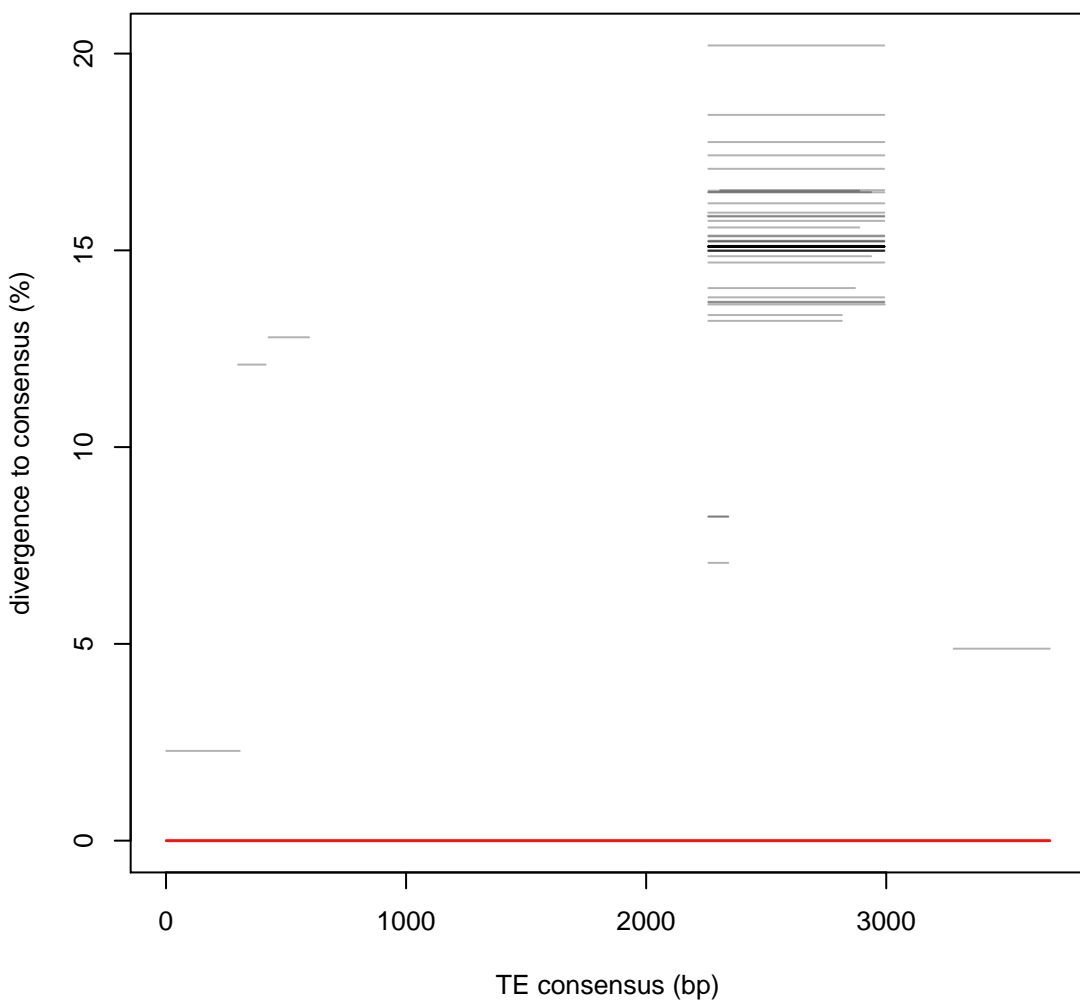
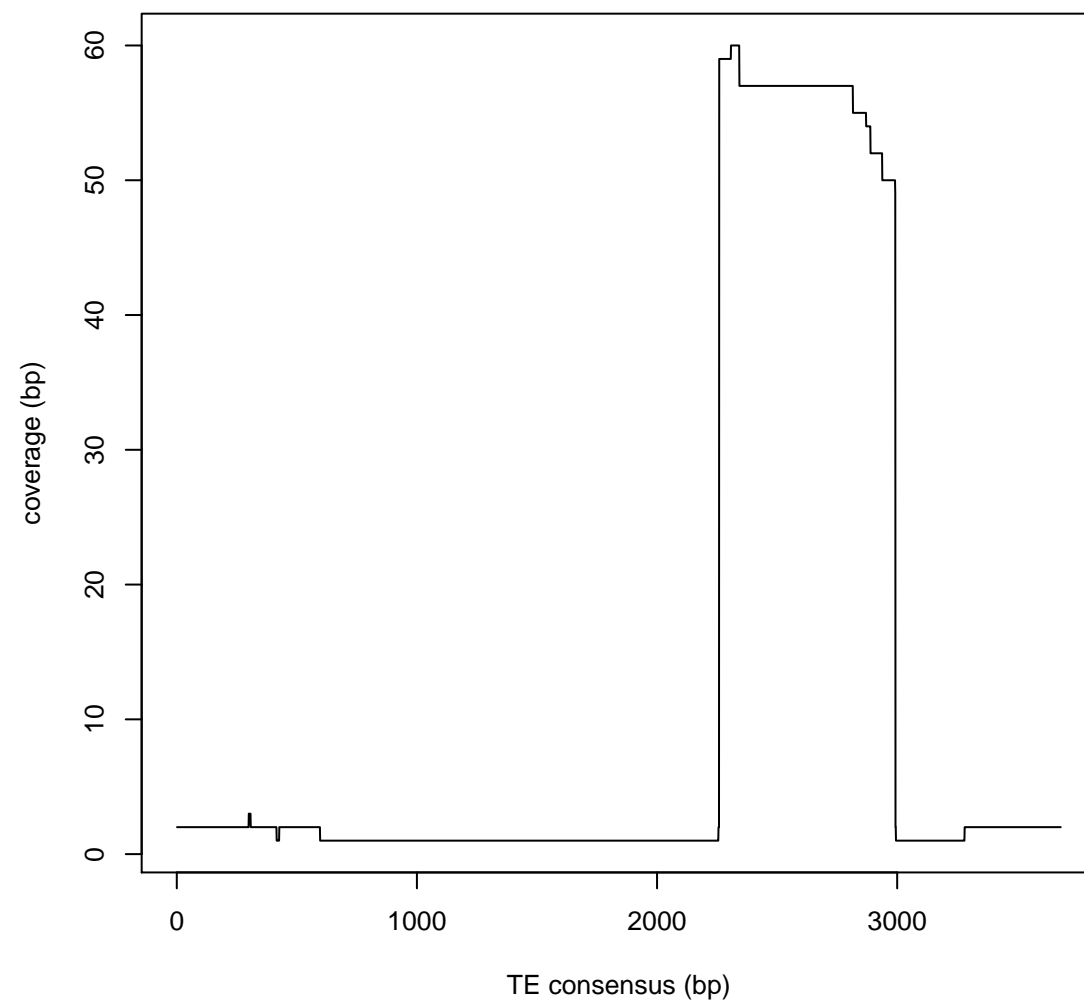


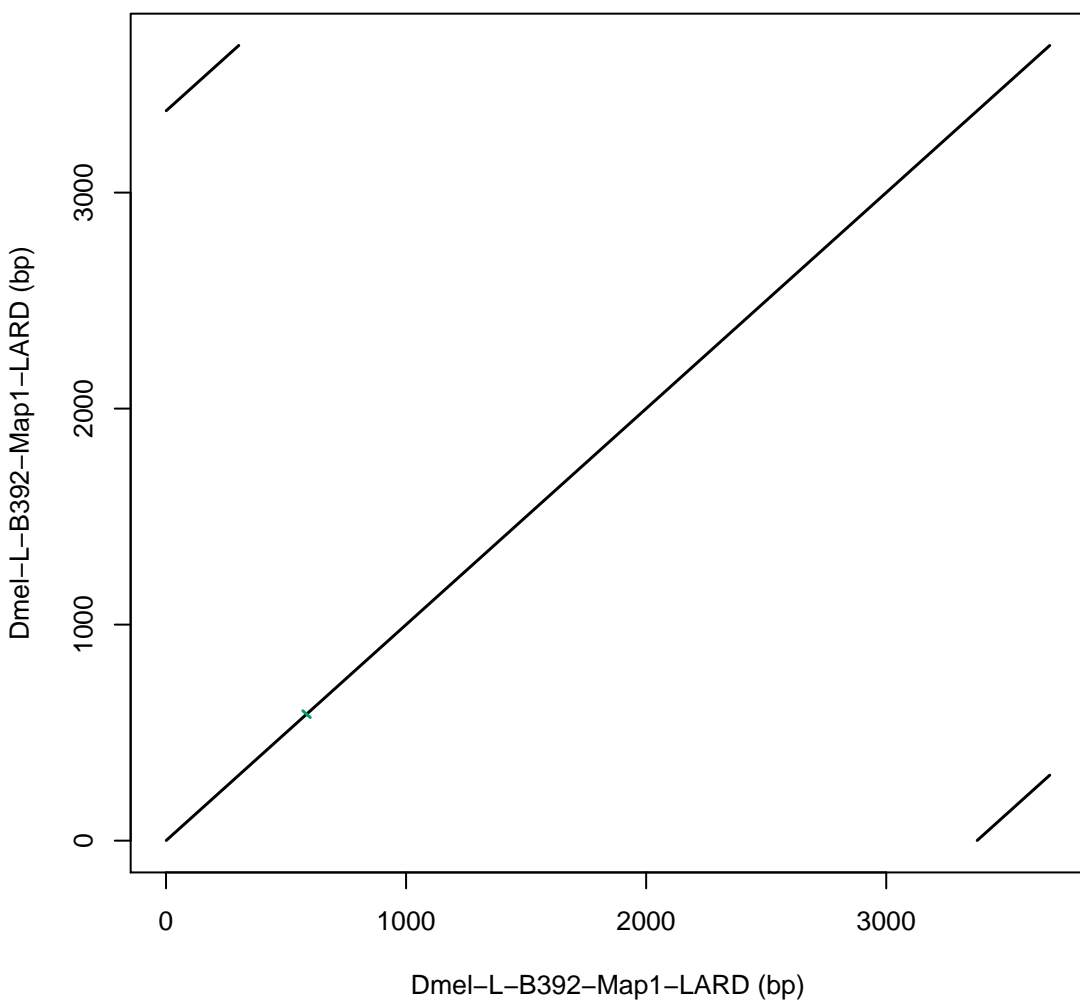
TE: Dmel-L-B392-Map1-LARD
consensus size: 3681bp; fragments: 64; full length: 1 (≥ 3312.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

