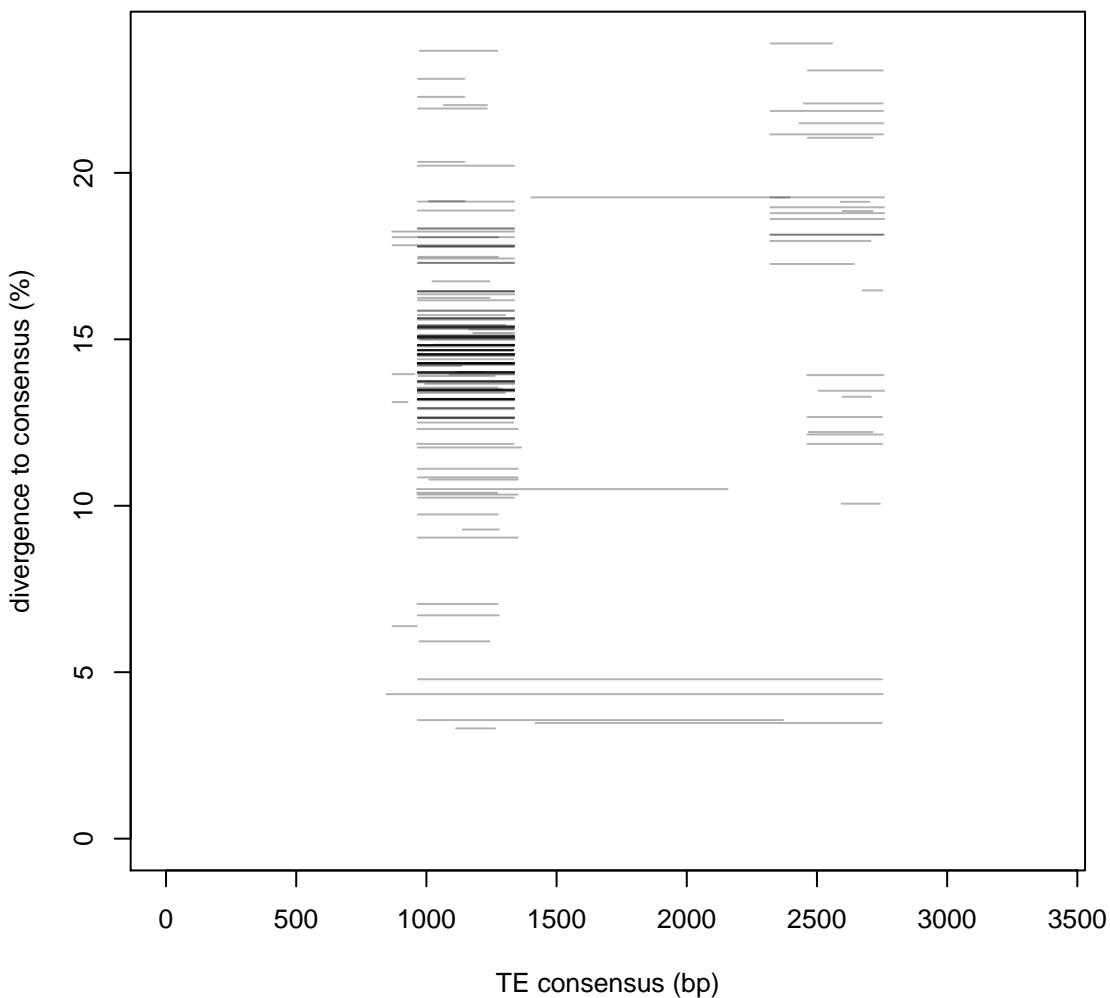
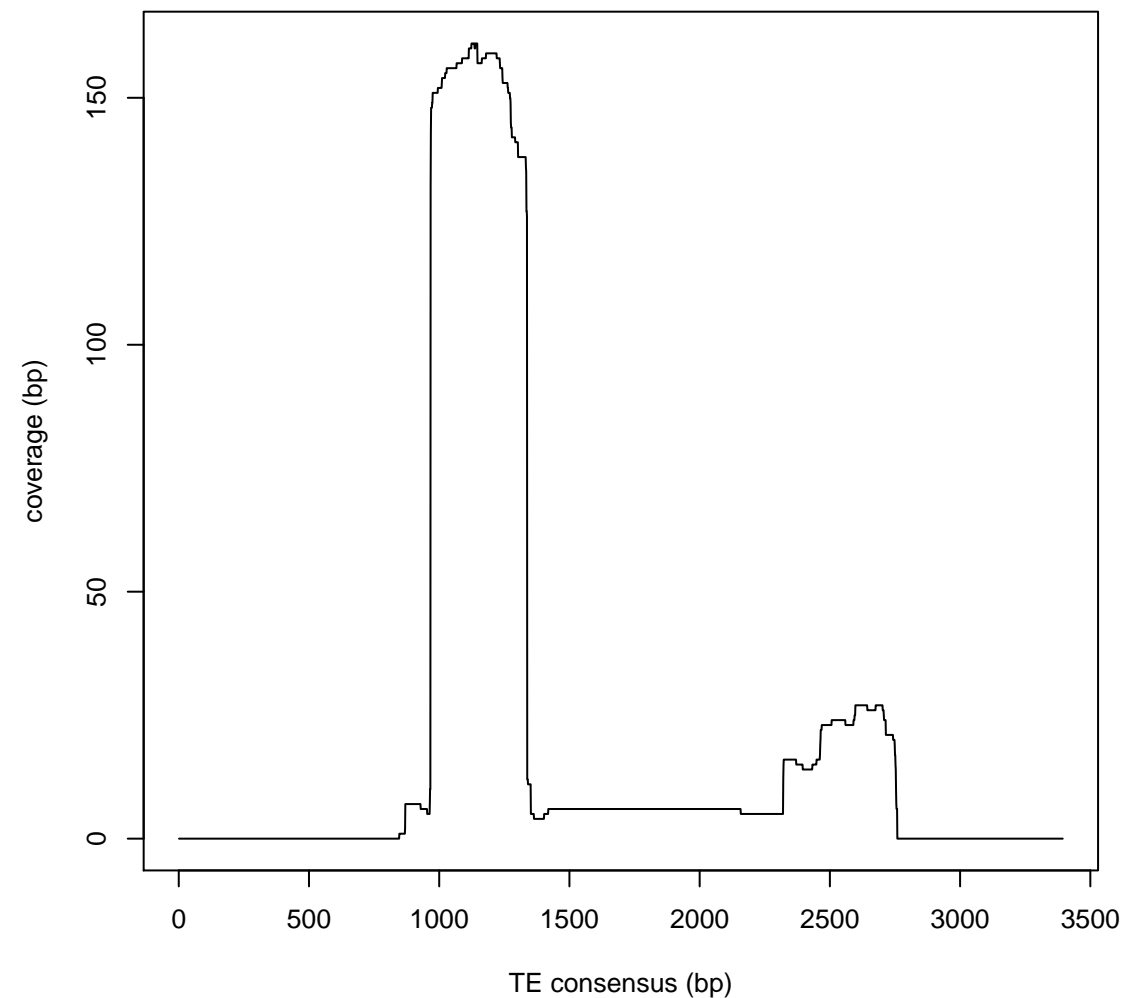


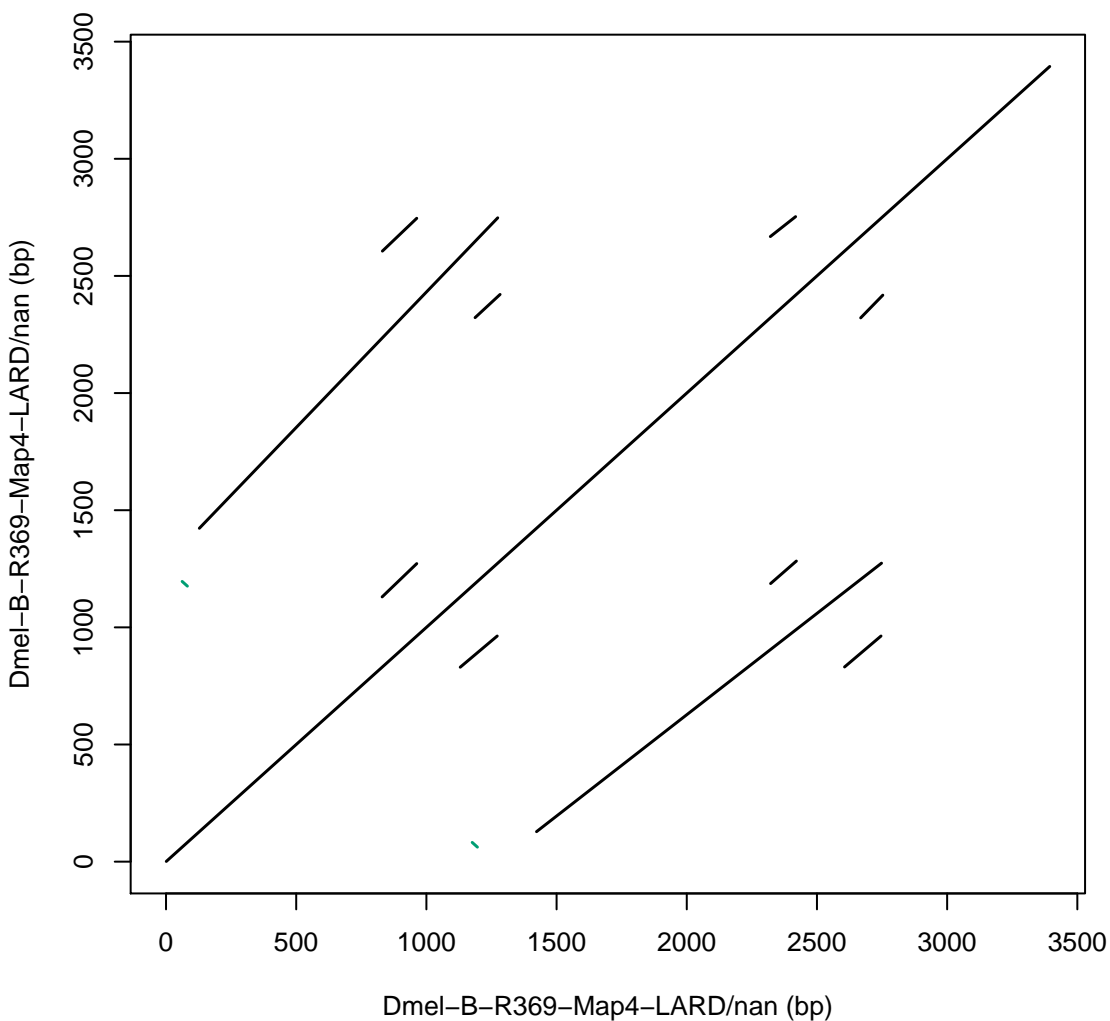
TE: Dmel-B-R369-Map4-LARD/nan
consensus size: 3394bp; fragments: 195; full length: 0 (≥ 3054.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

