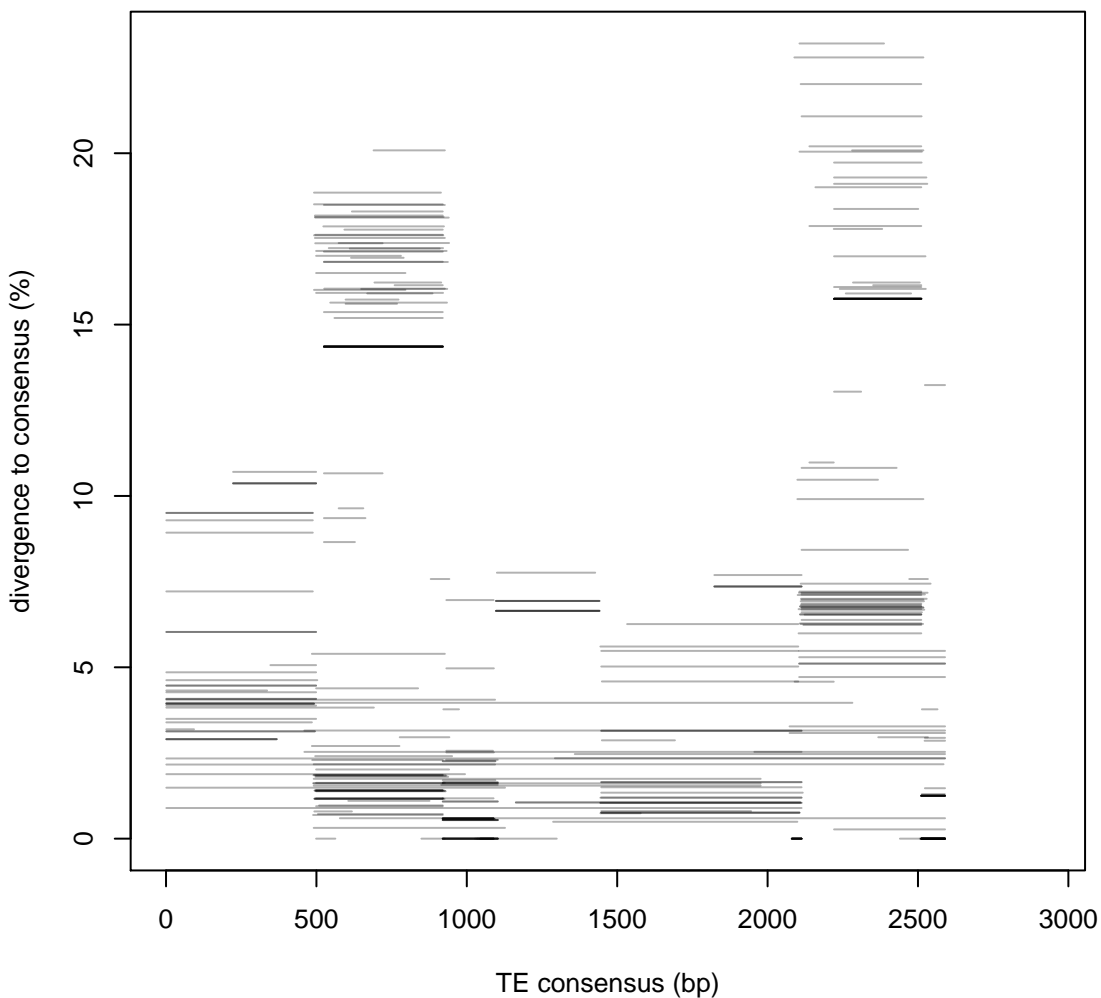
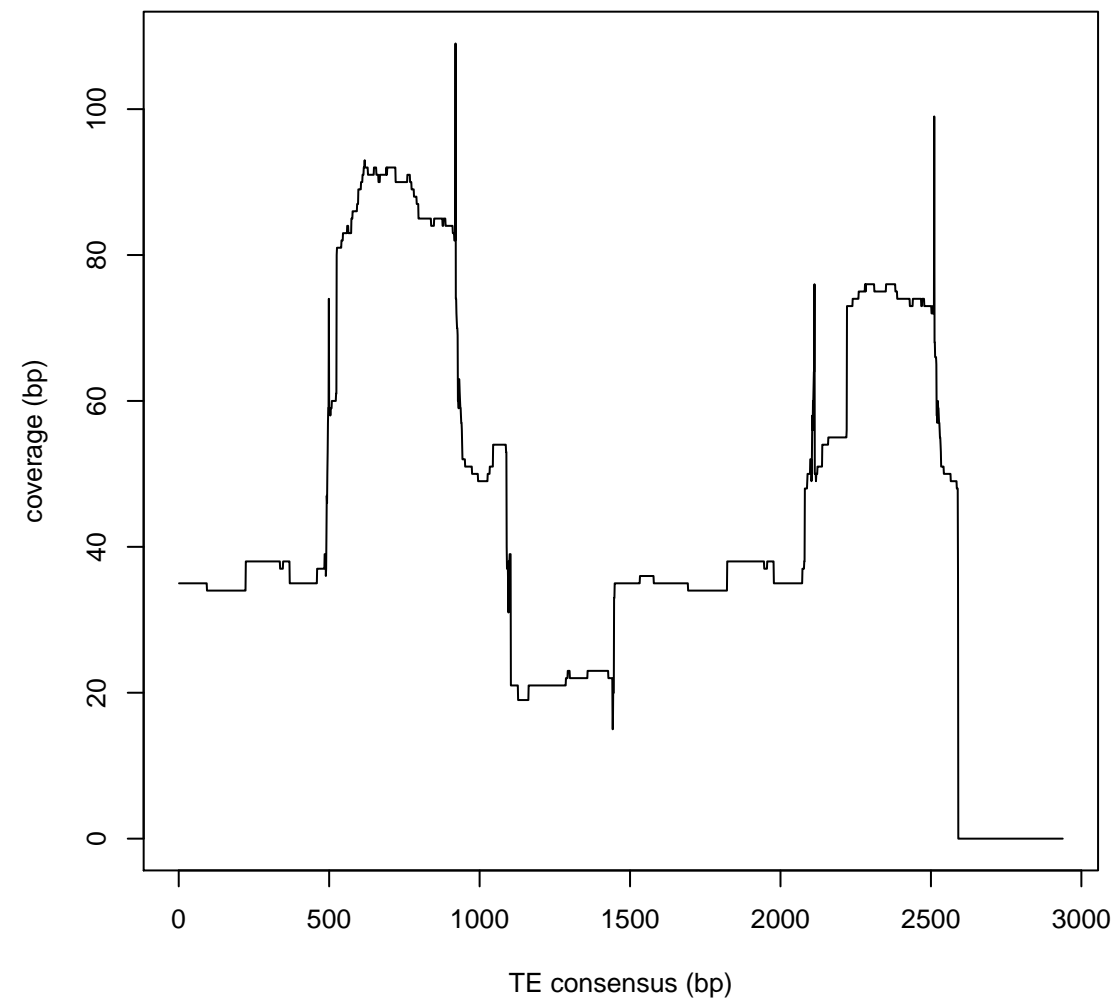


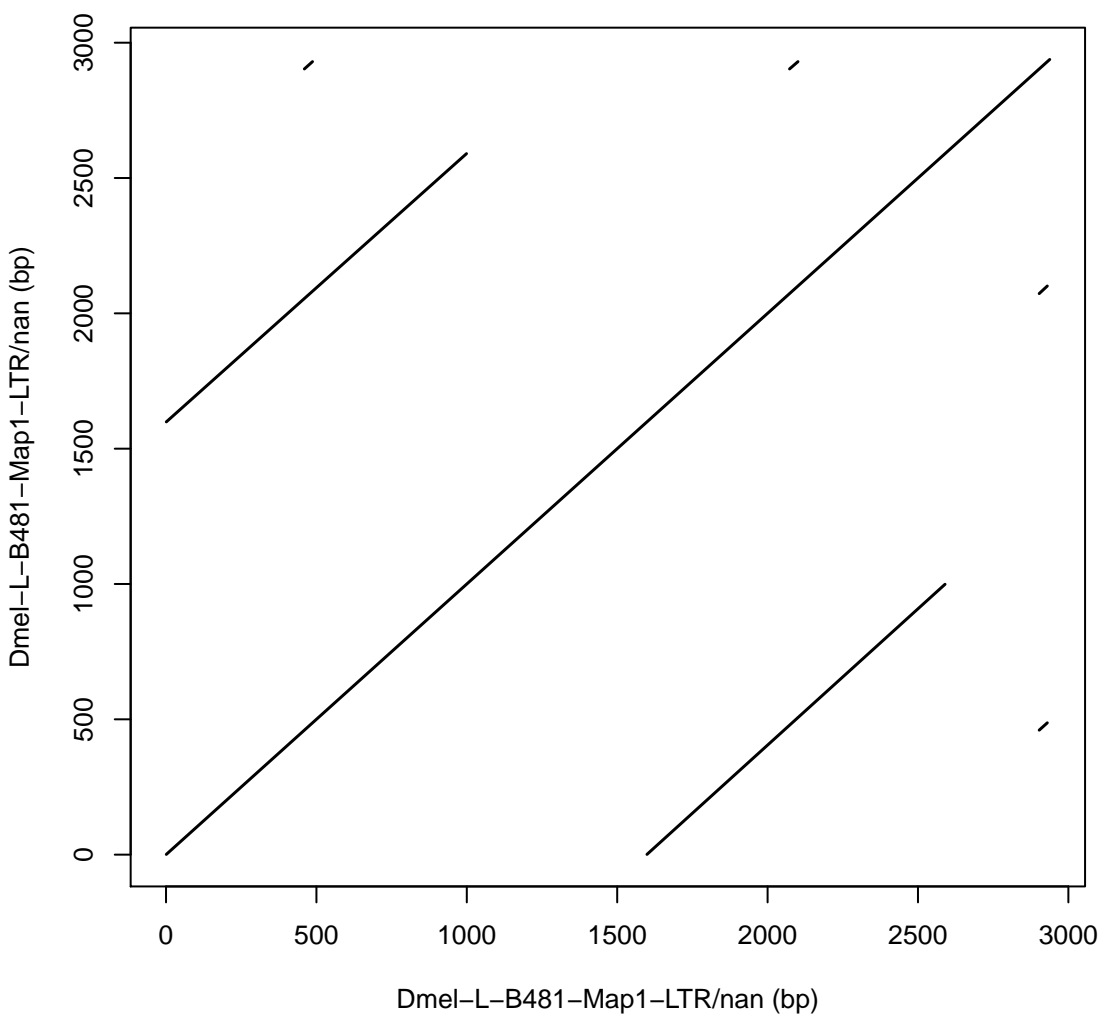
TE: Dmel-L-B481-Map1-LTR/nan  
consensus size: 2938bp; fragments: 335; full length: 0 (>=2644.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

