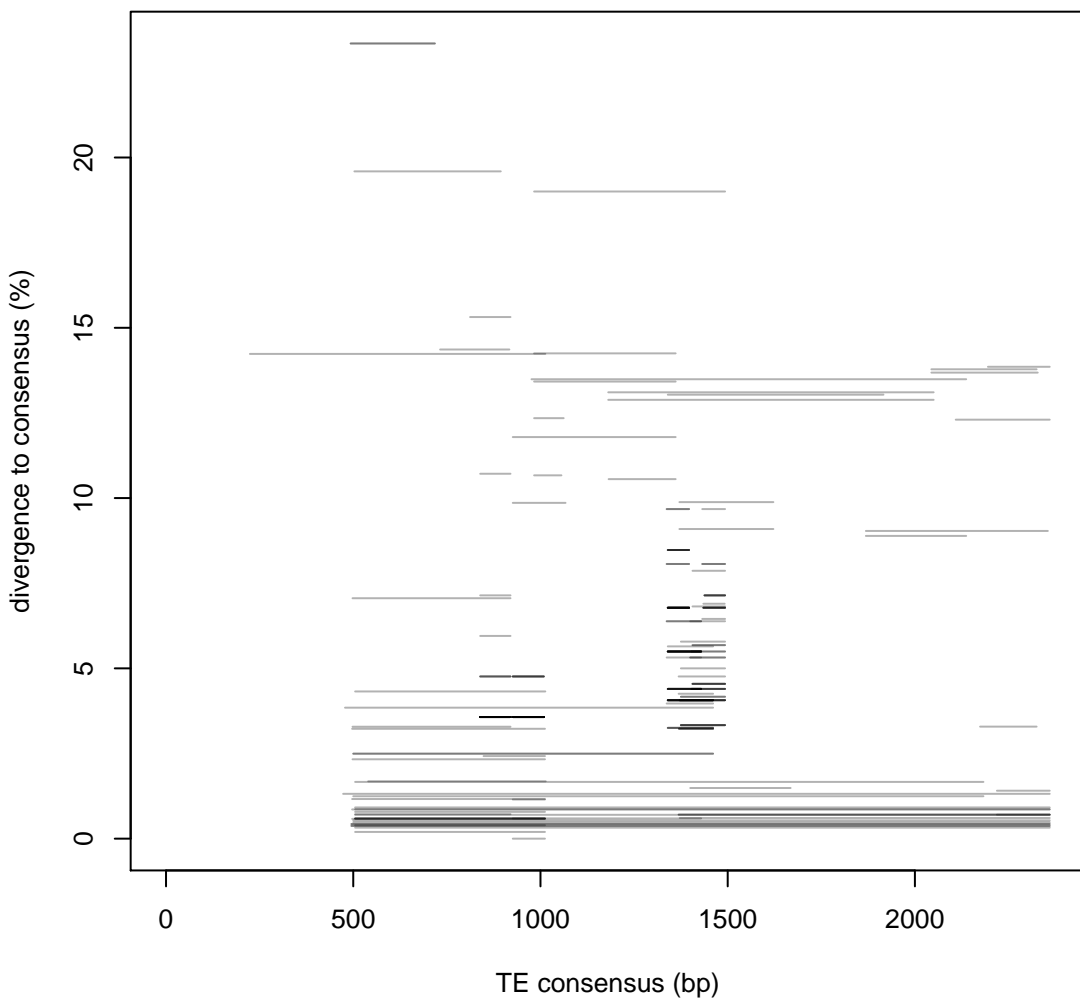
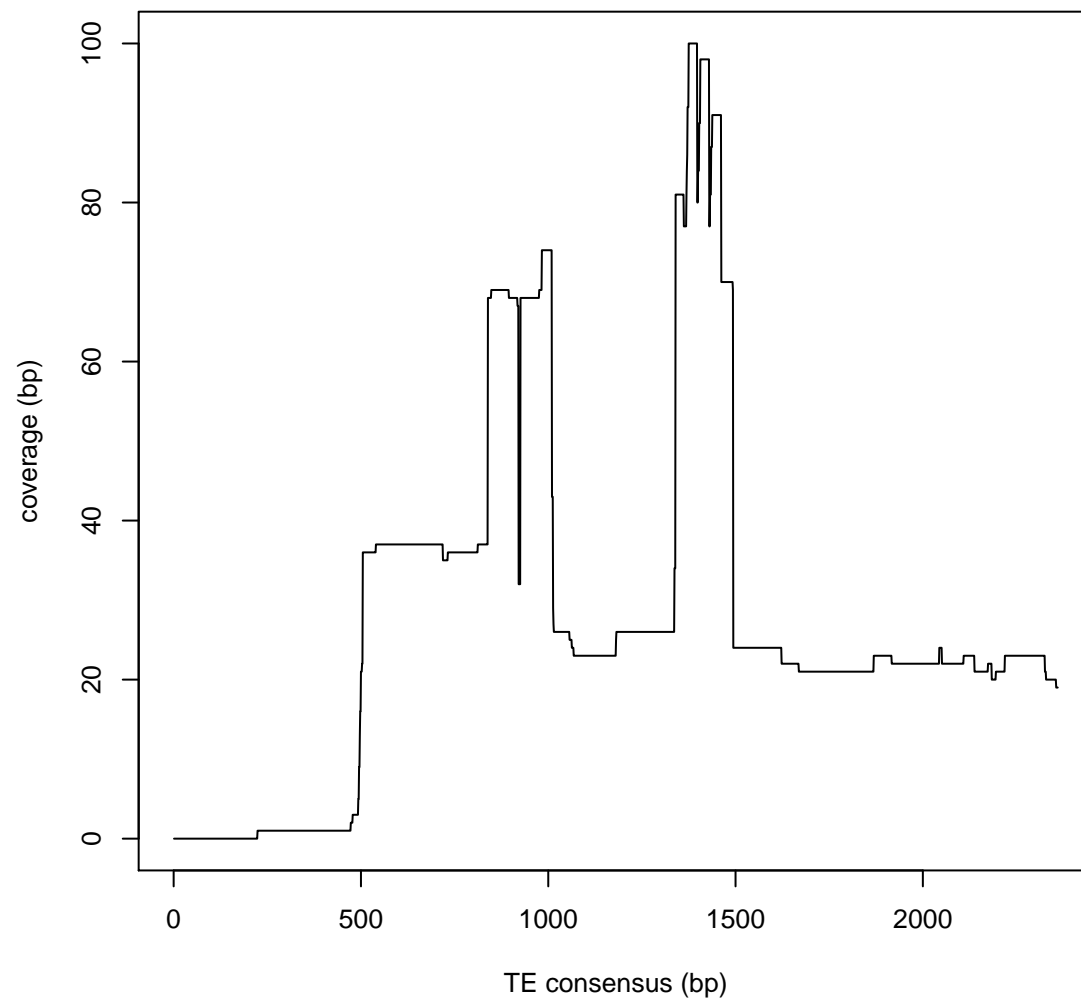


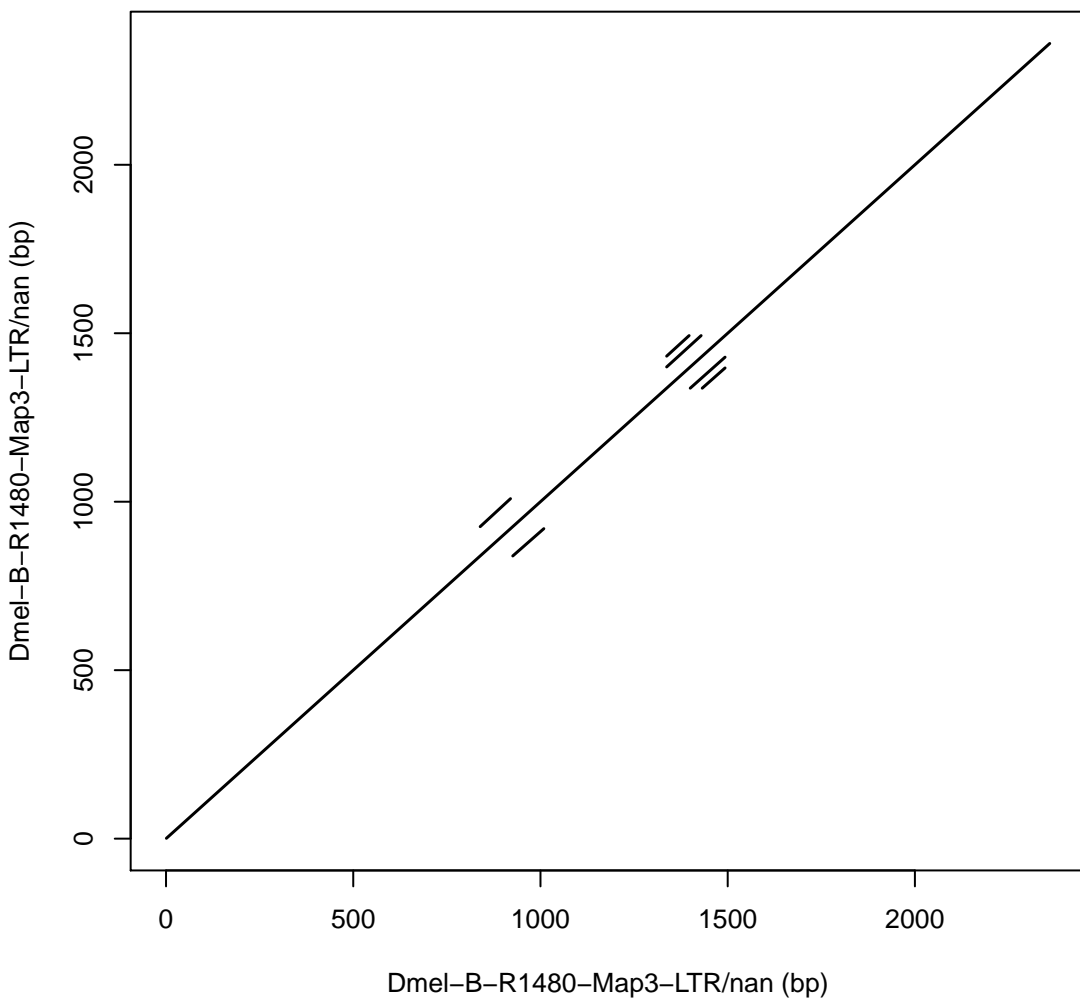
TE: Dmel-B-R1480-Map3-LTR/nan  
consensus size: 2360bp; fragments: 235; full length: 0 (>=2124bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

