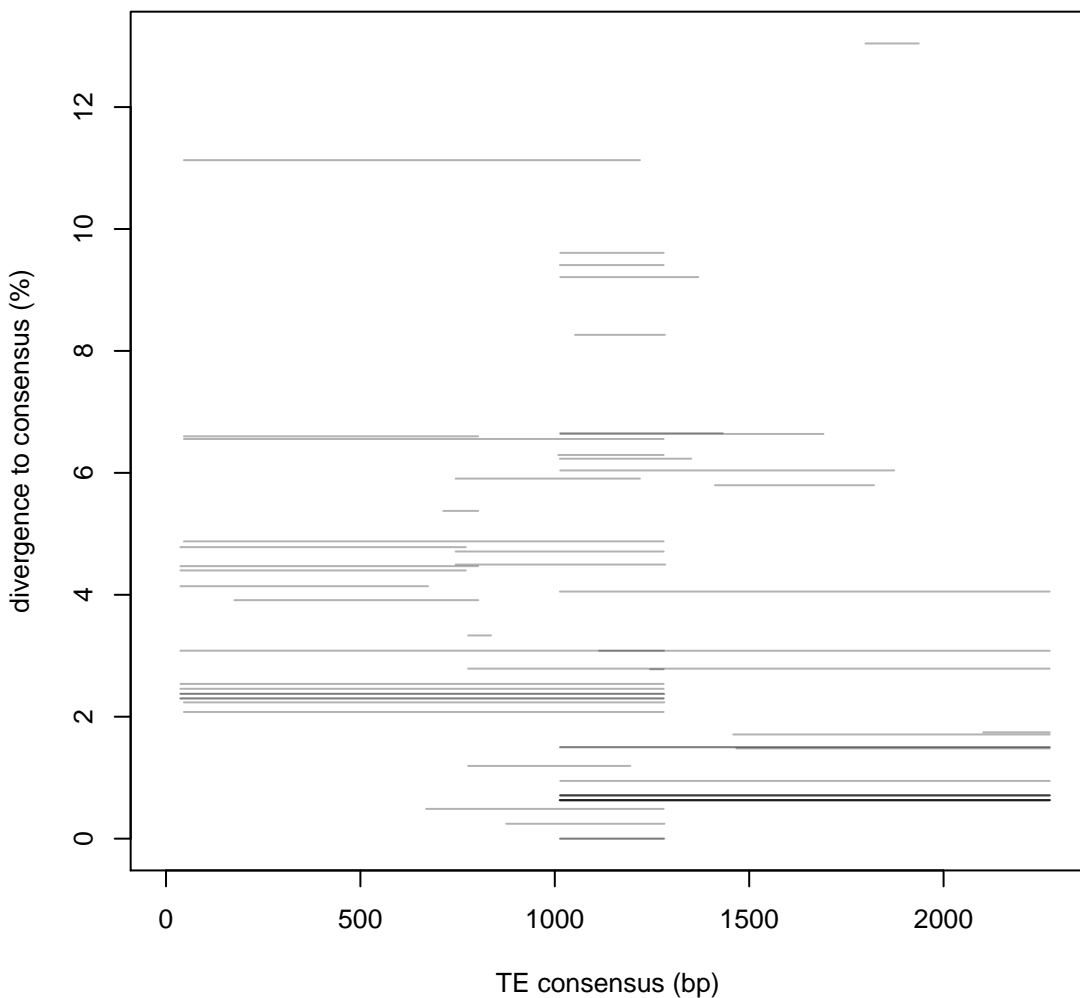
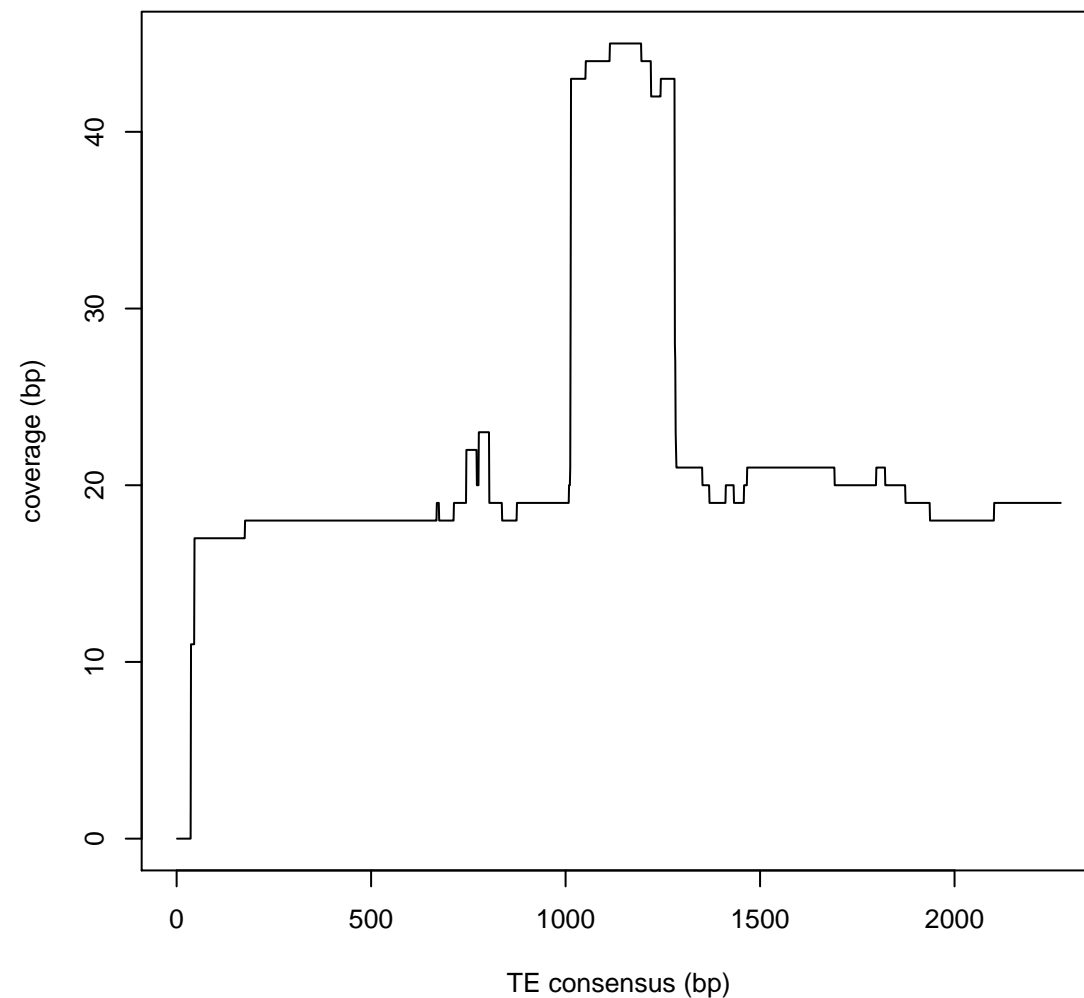


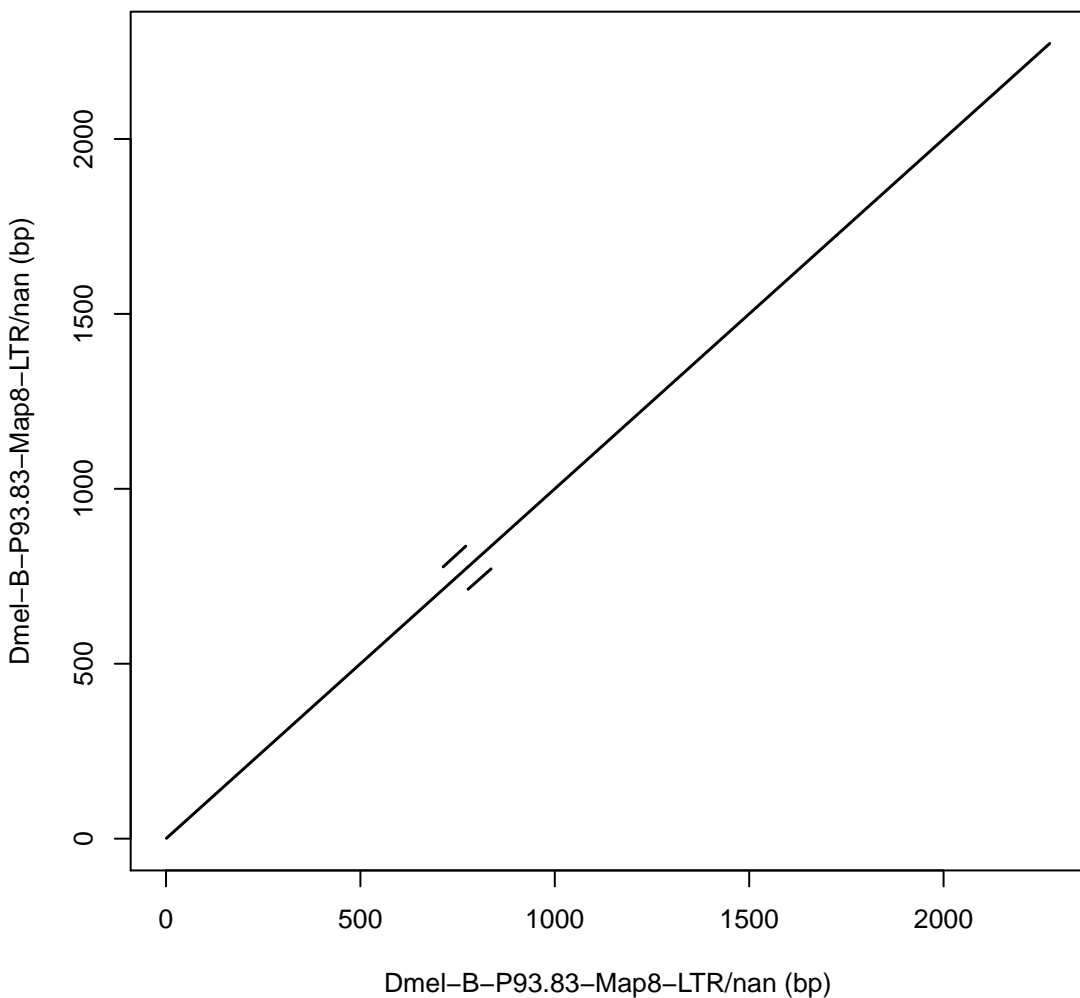
TE: Dmel-B-P93.83-Map8-LTR/nan  
consensus size: 2273bp; fragments: 59; full length: 0 (>=2045.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

