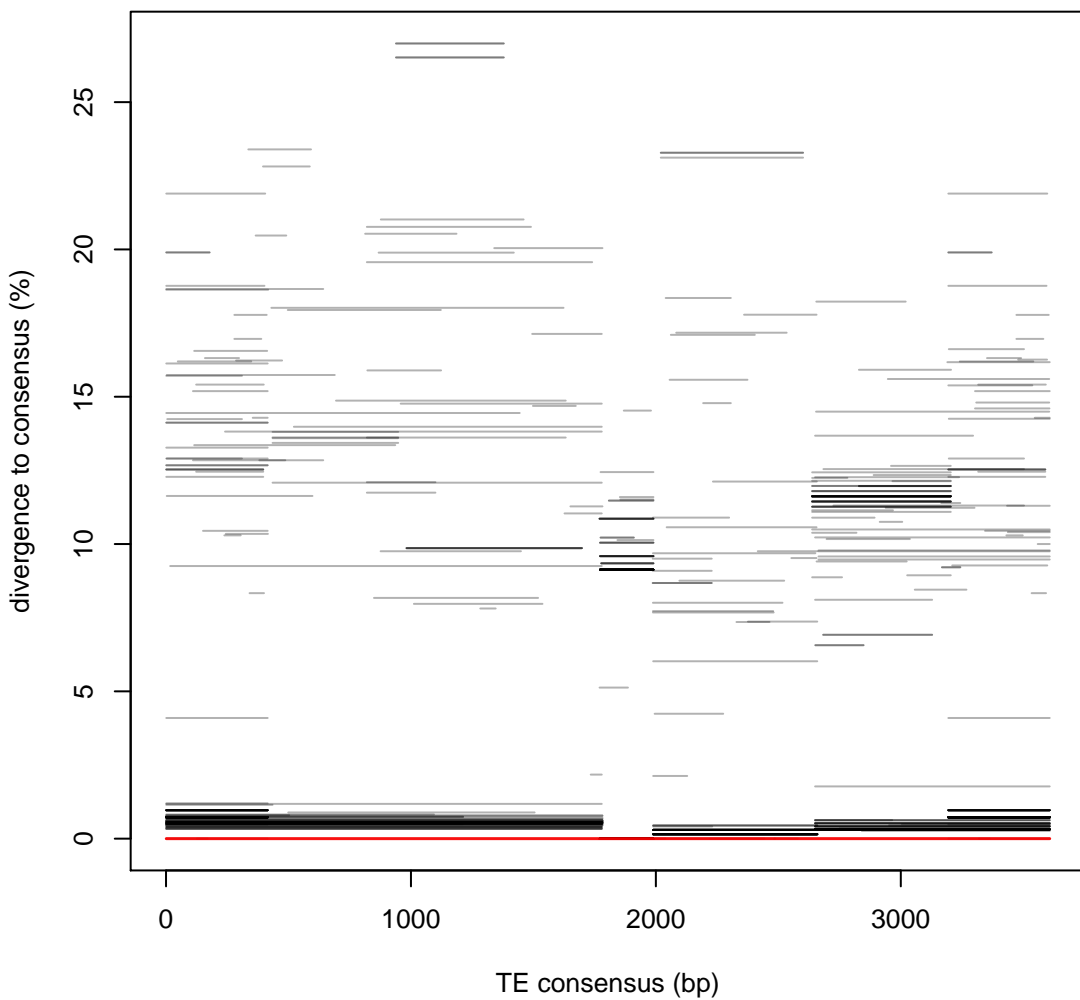
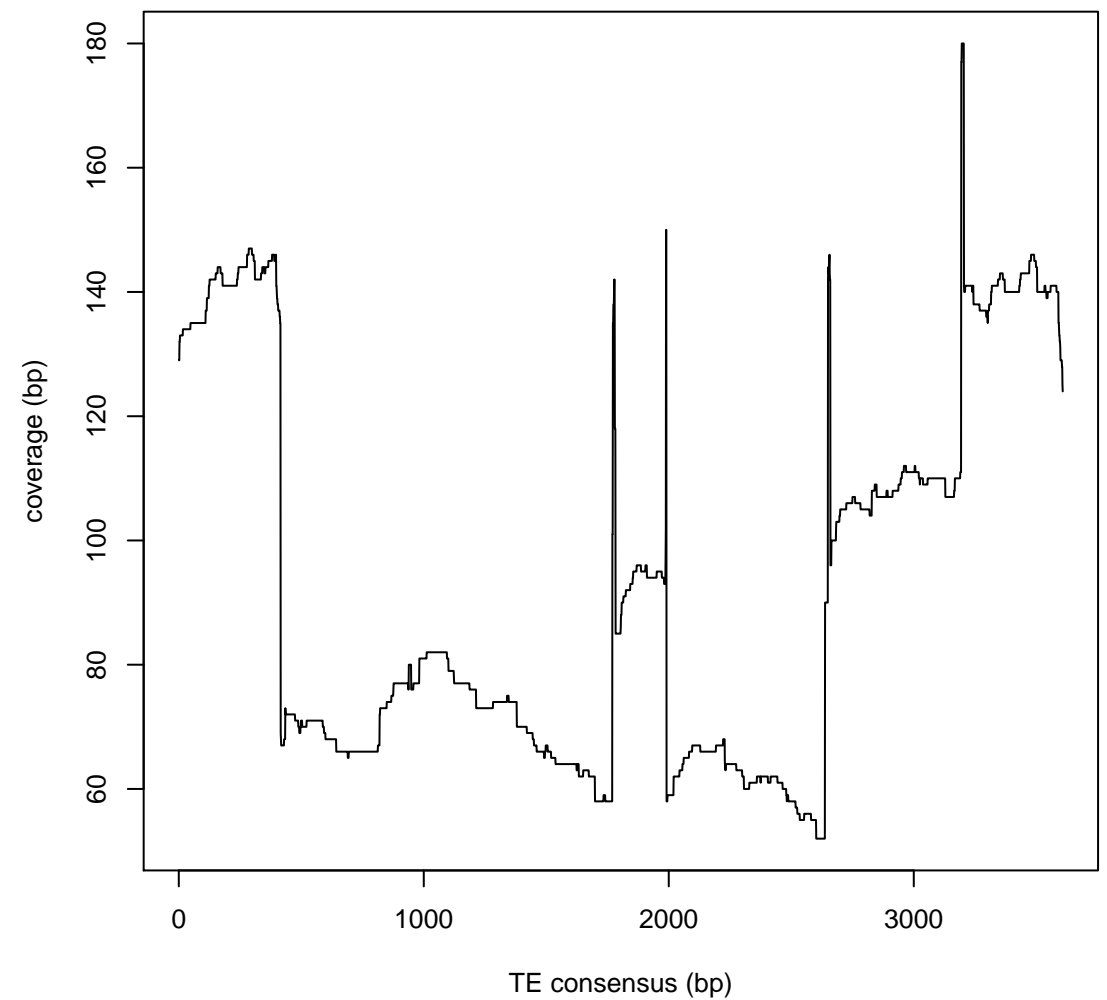


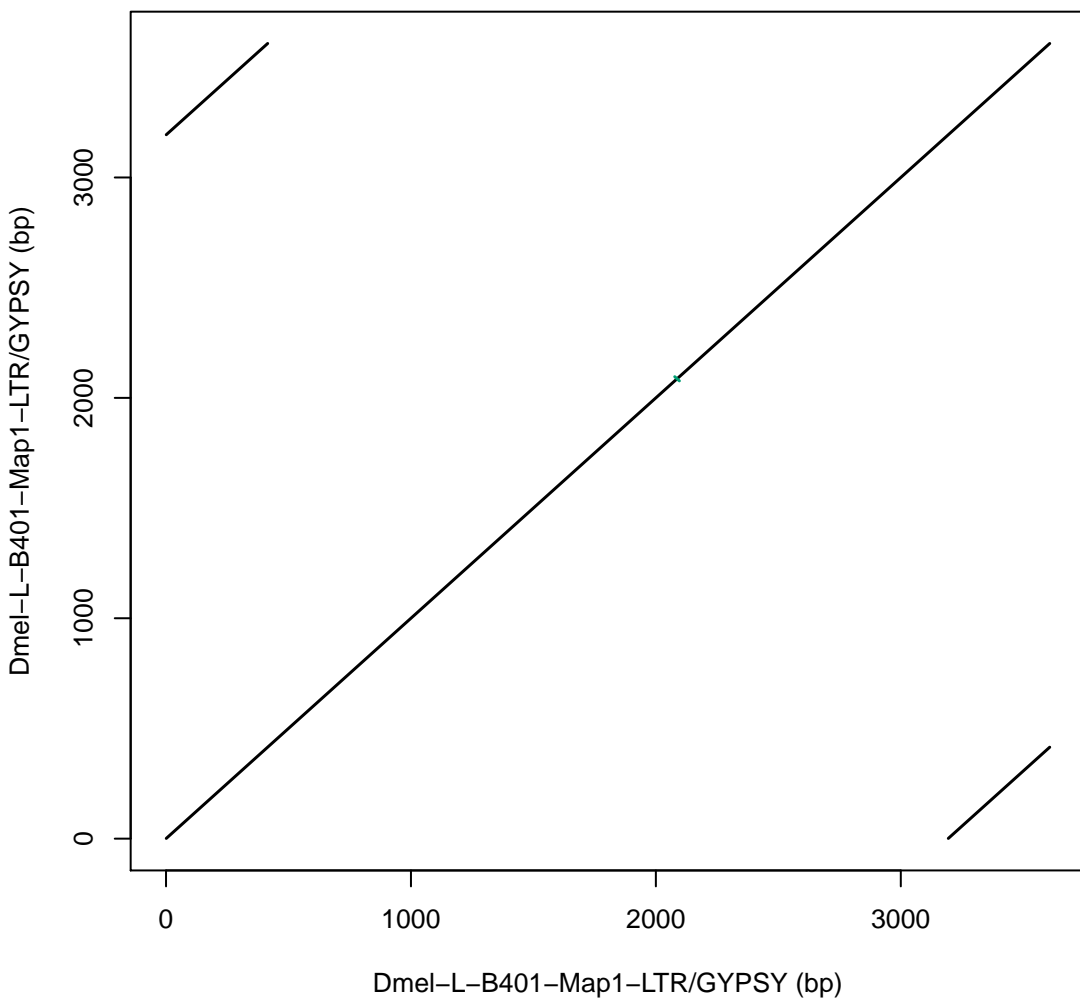
TE: Dmel-L-B401-Map1-LTR/GYPY
consensus size: 3608bp; fragments: 580; full length: 1 (>=3247.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

