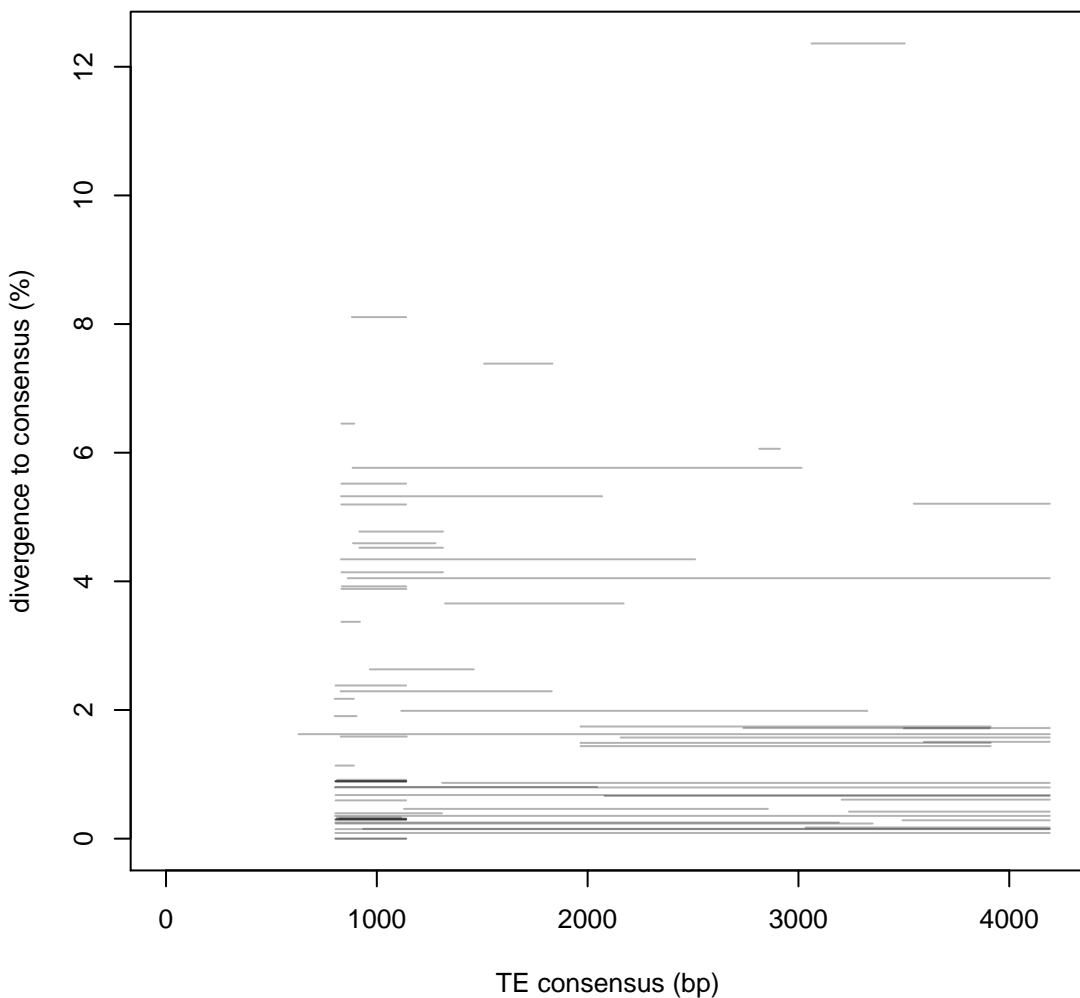
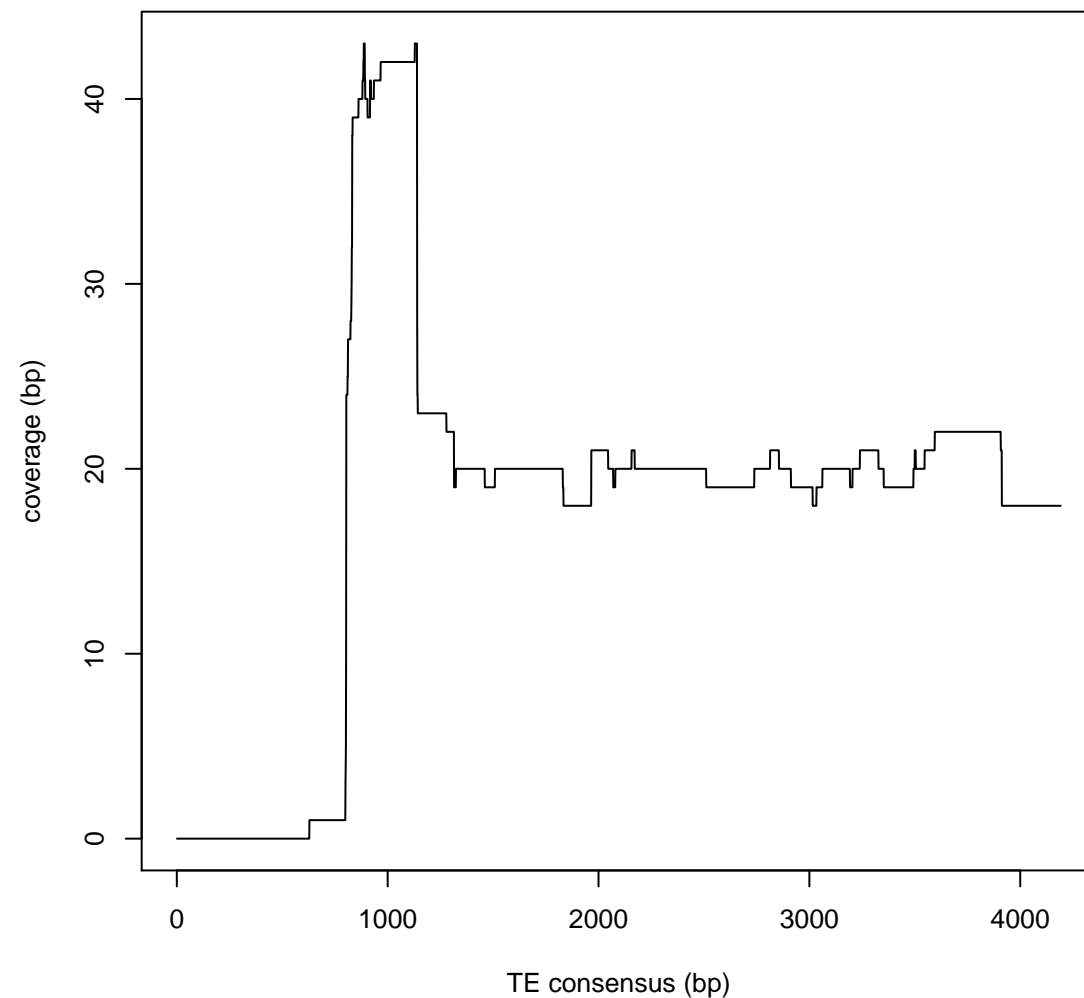


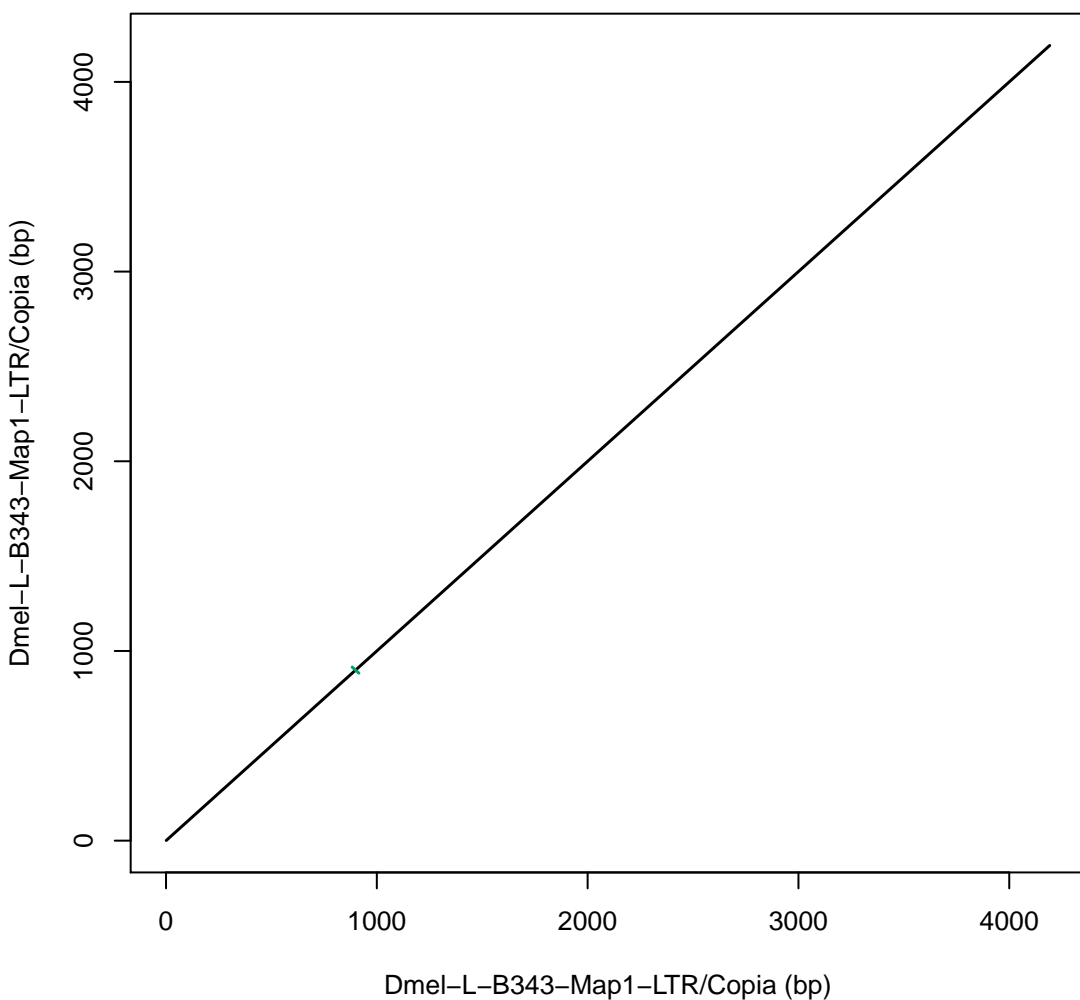
TE: Dmel-L-B343-Map1-LTR/Copia
consensus size: 4192bp; fragments: 67; full length: 0 (≥ 3772.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

