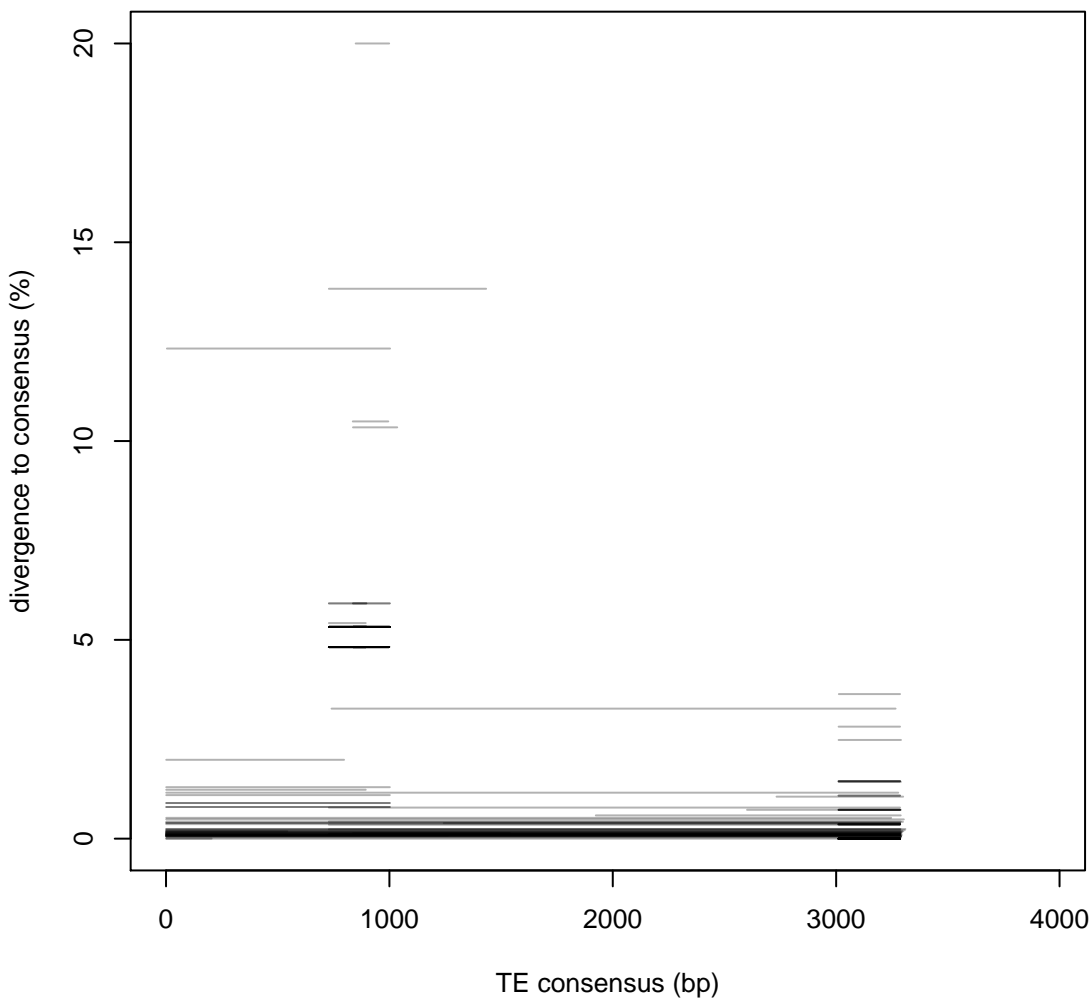
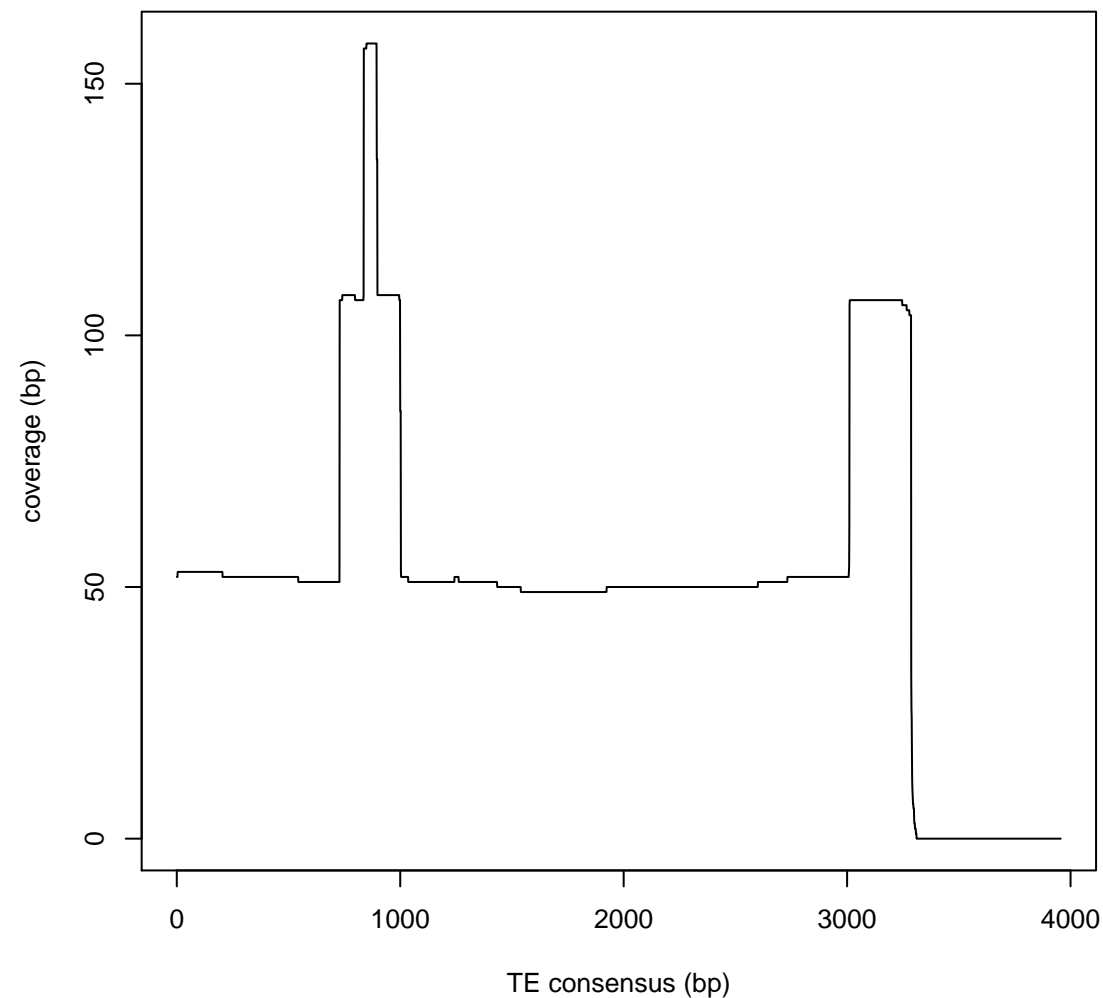


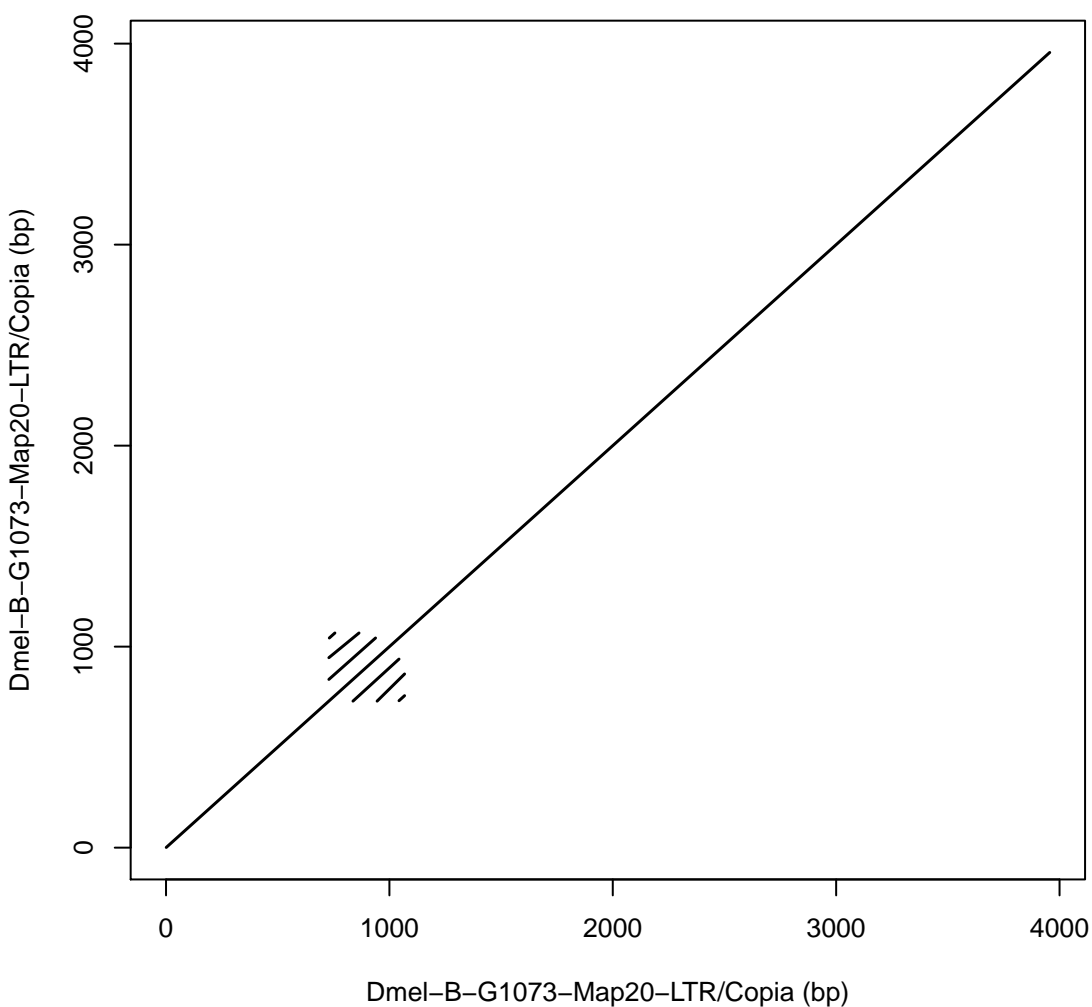
TE: Dmel-B-G1073-Map20-LTR/Copia
consensus size: 3956bp; fragments: 220; full length: 0 (>=3560.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

