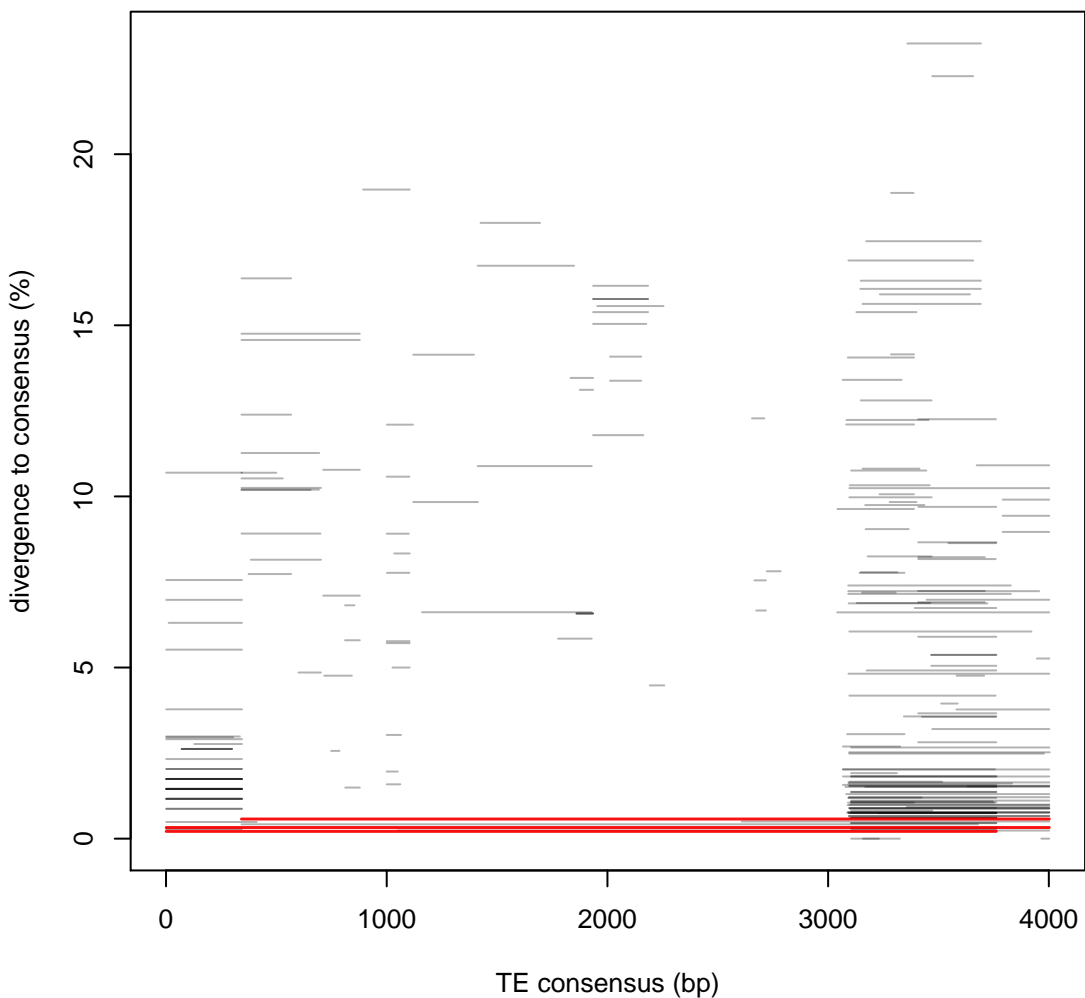
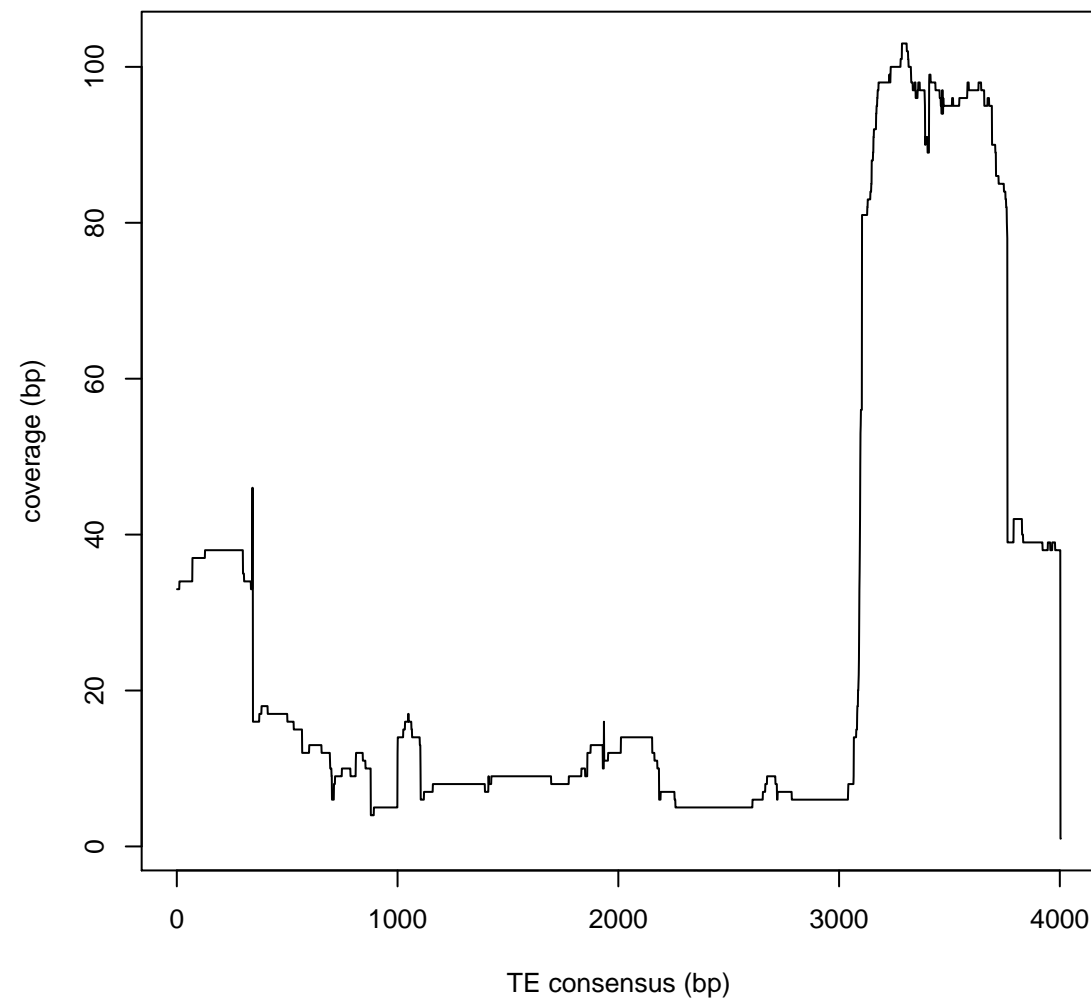


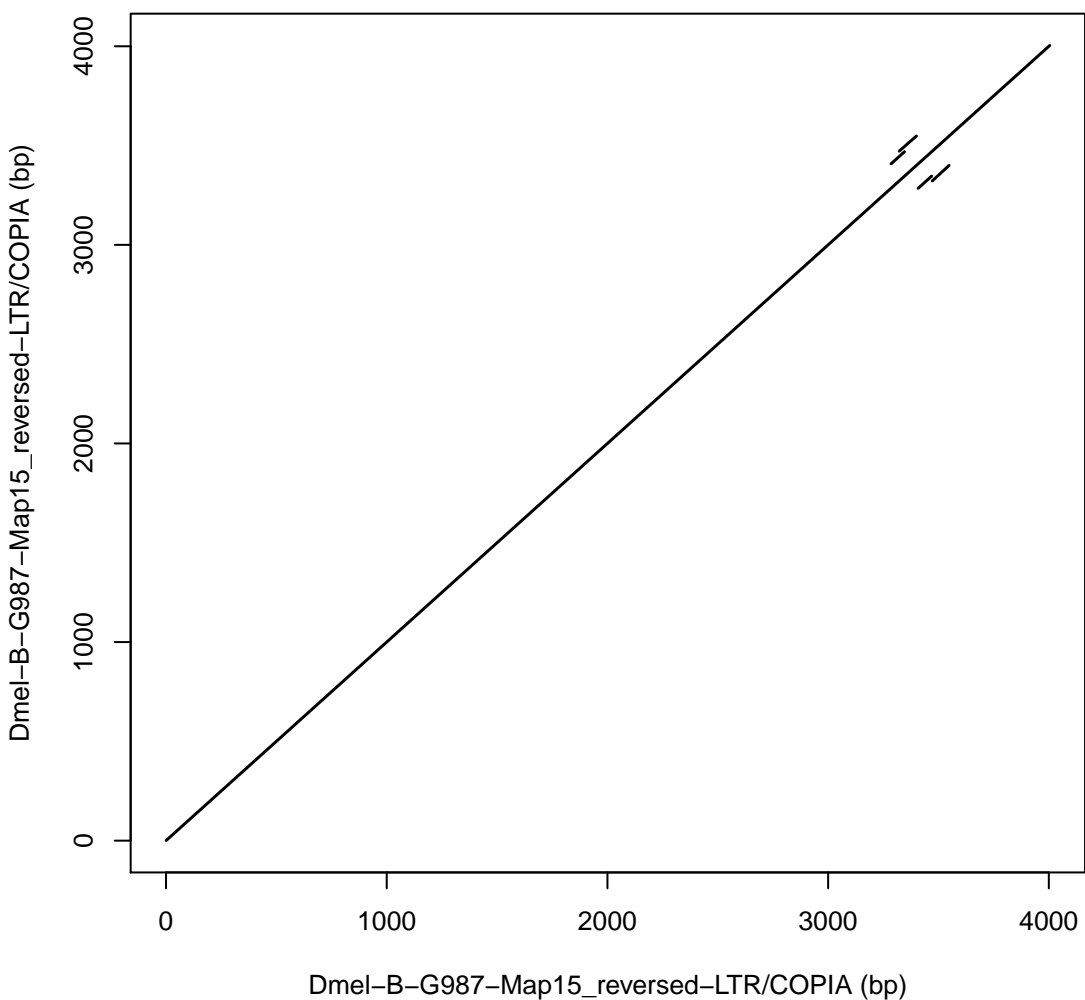
TE: Dmel-B-G987-Map15_reversed-LTR/COPIA
consensus size: 4004bp; fragments: 231; full length: 3 (>=3603.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

