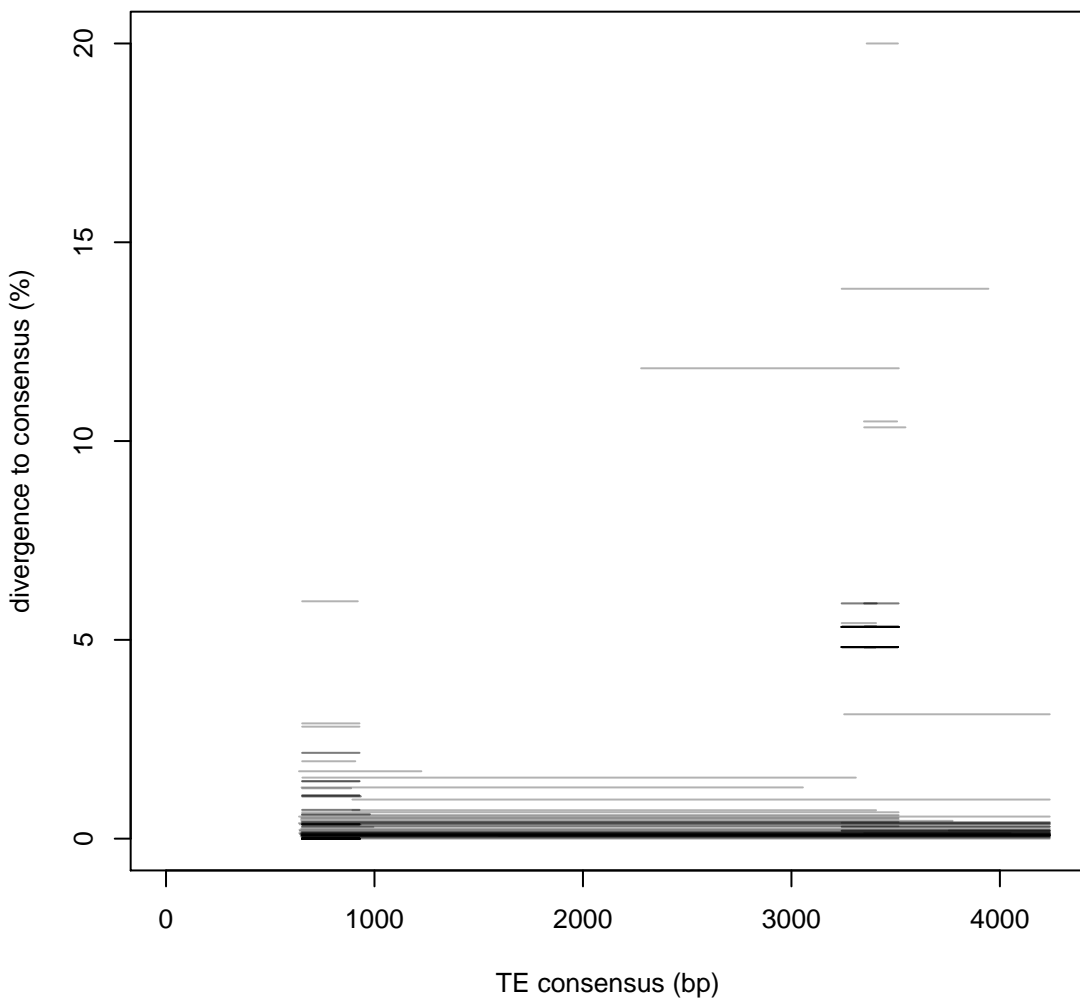
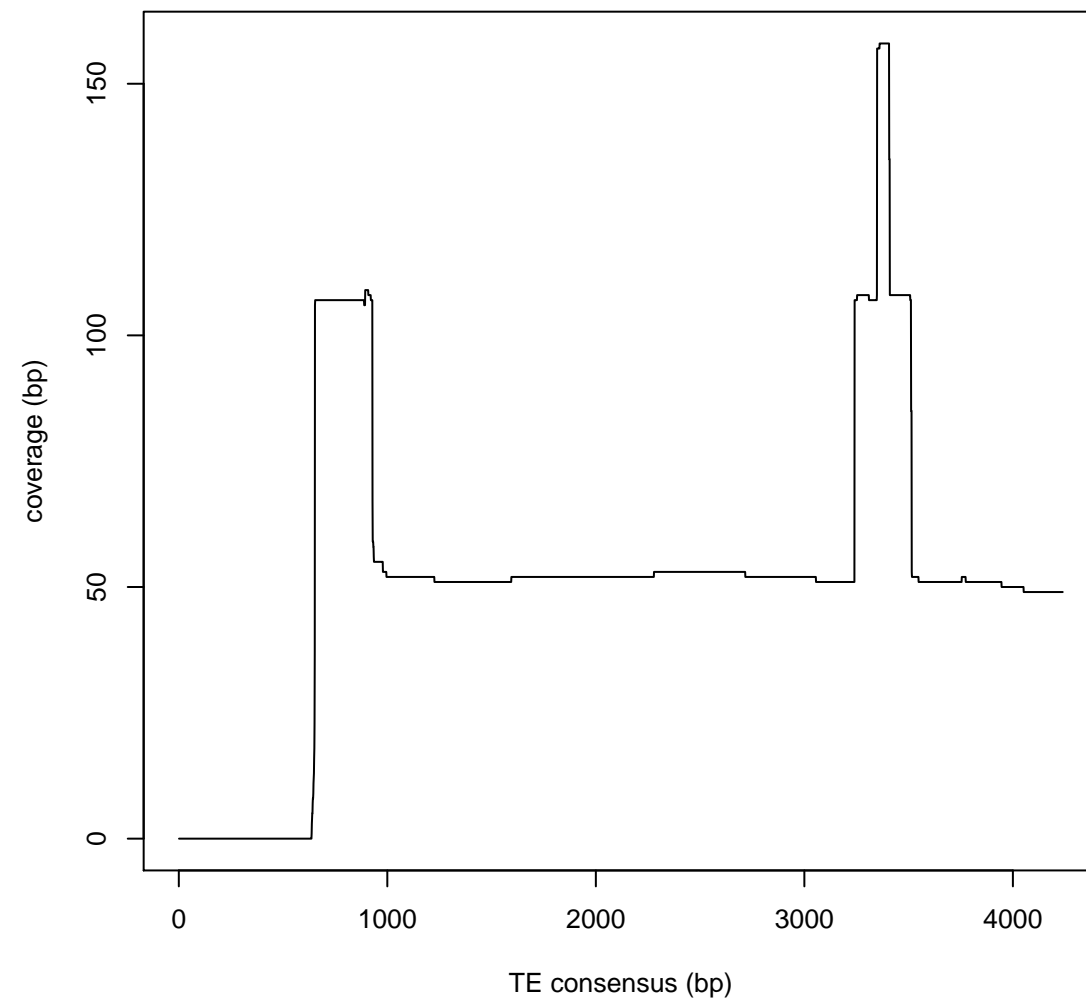


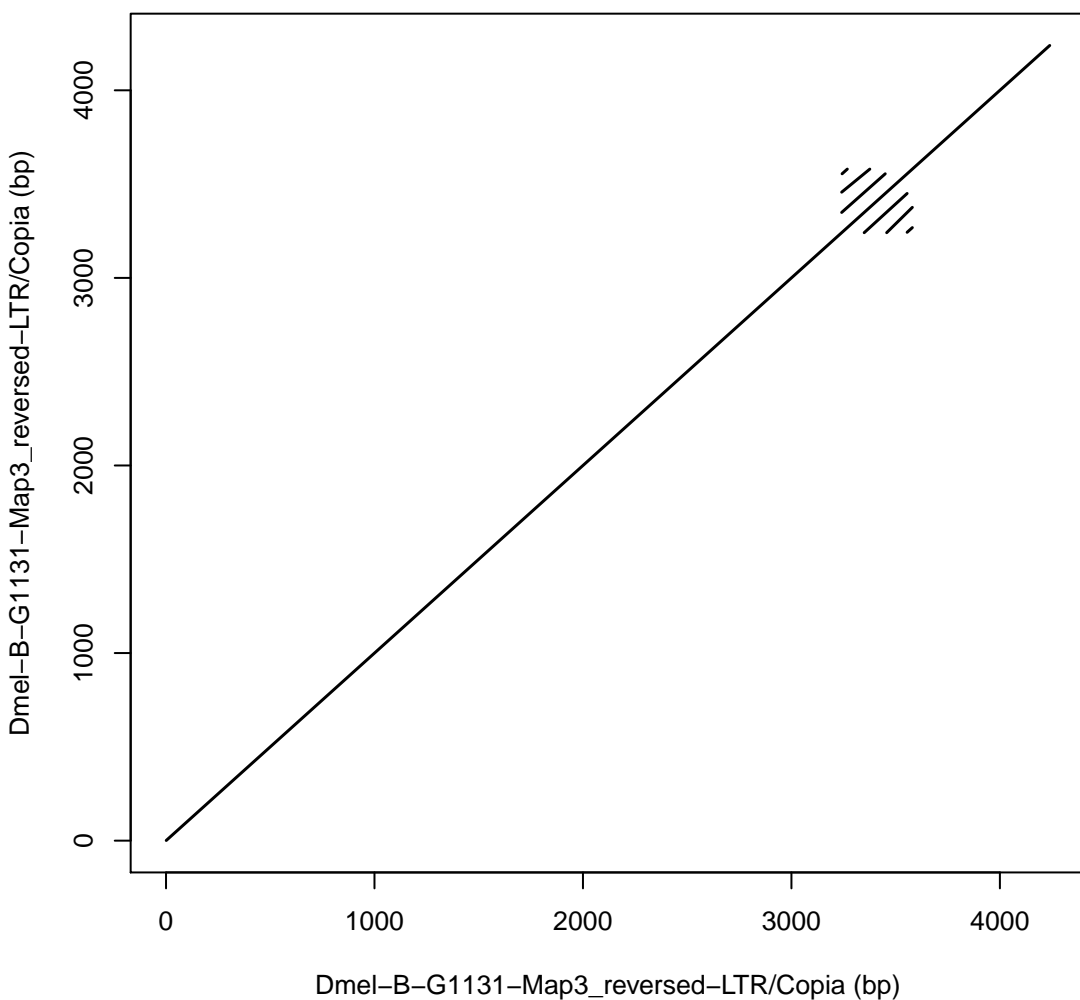
TE: Dmel-B-G1131-Map3_reversed-LTR/Copia
consensus size: 4239bp; fragments: 221; full length: 0 (>=3815.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

