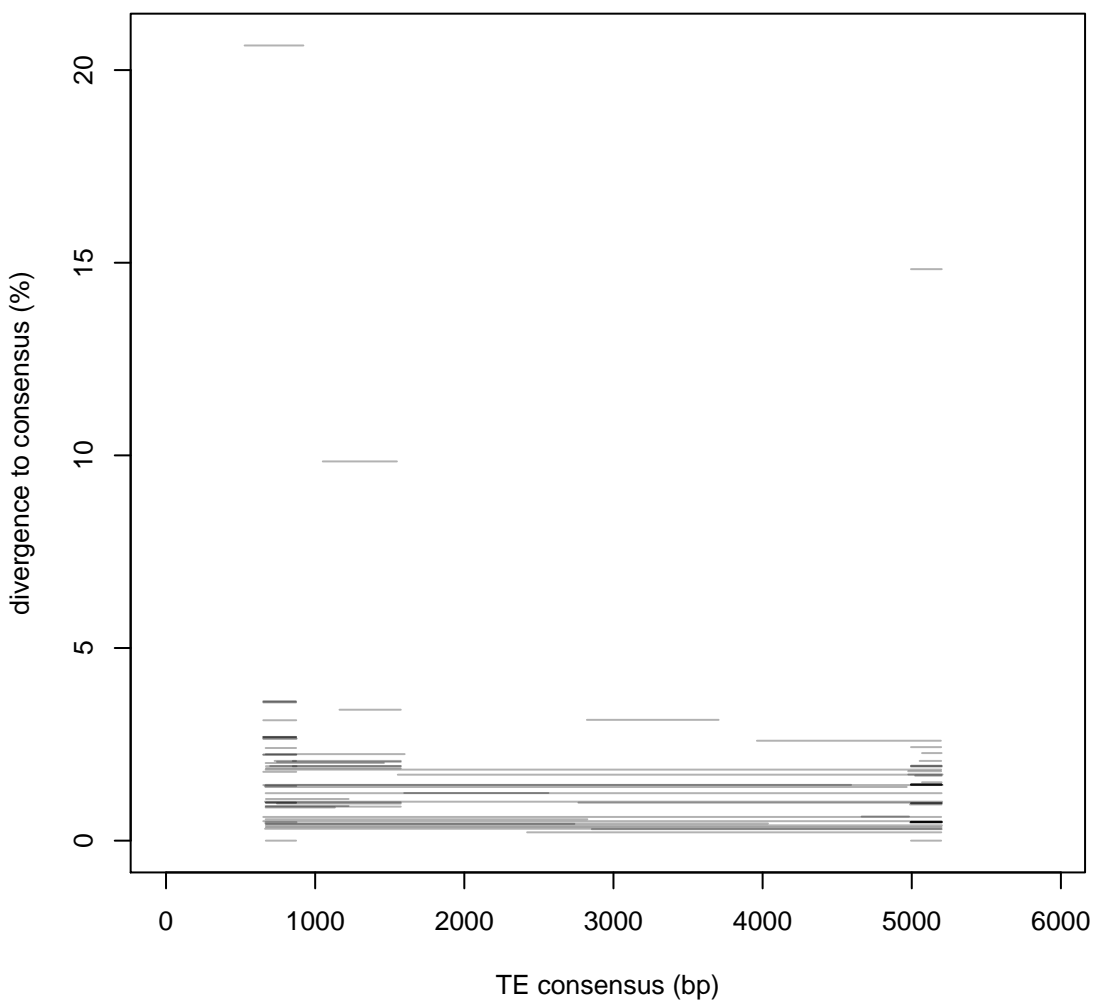
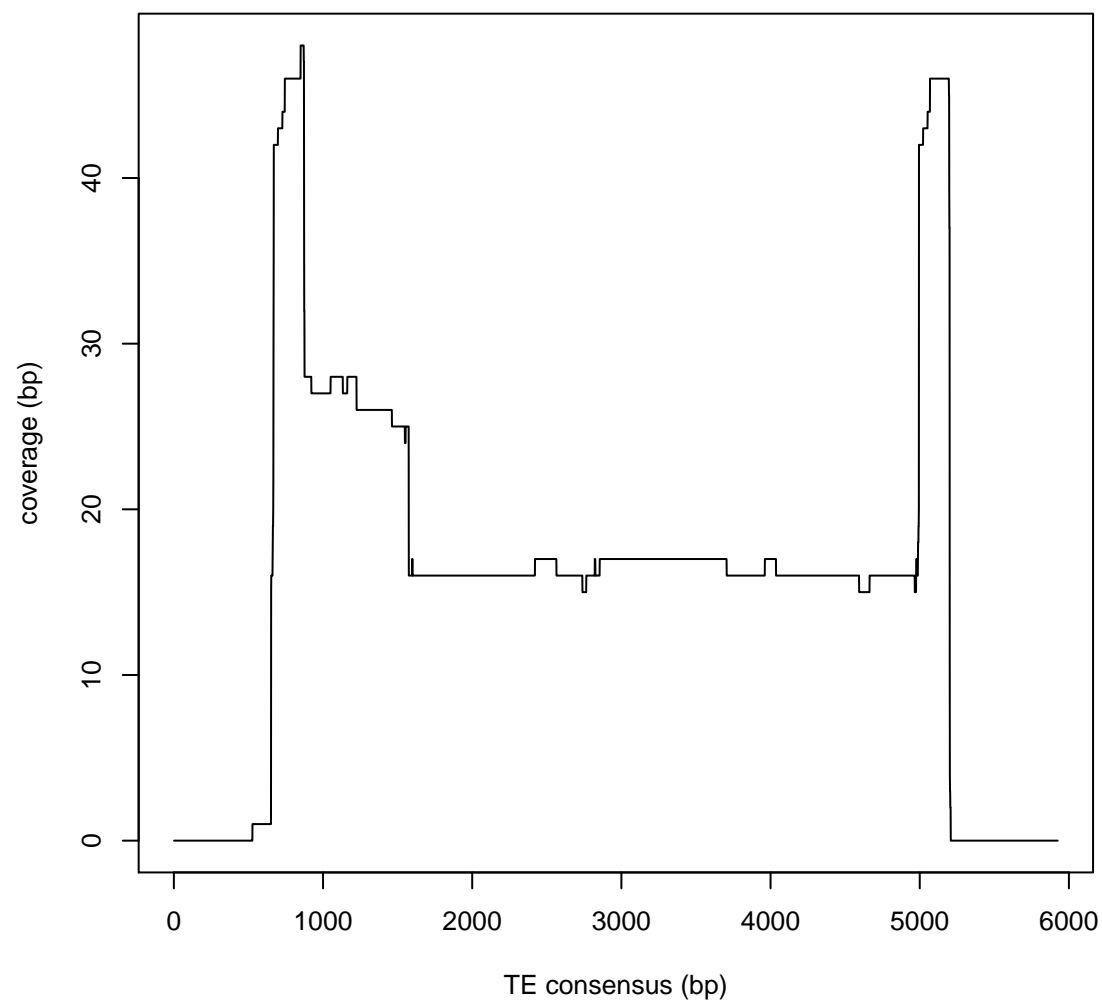


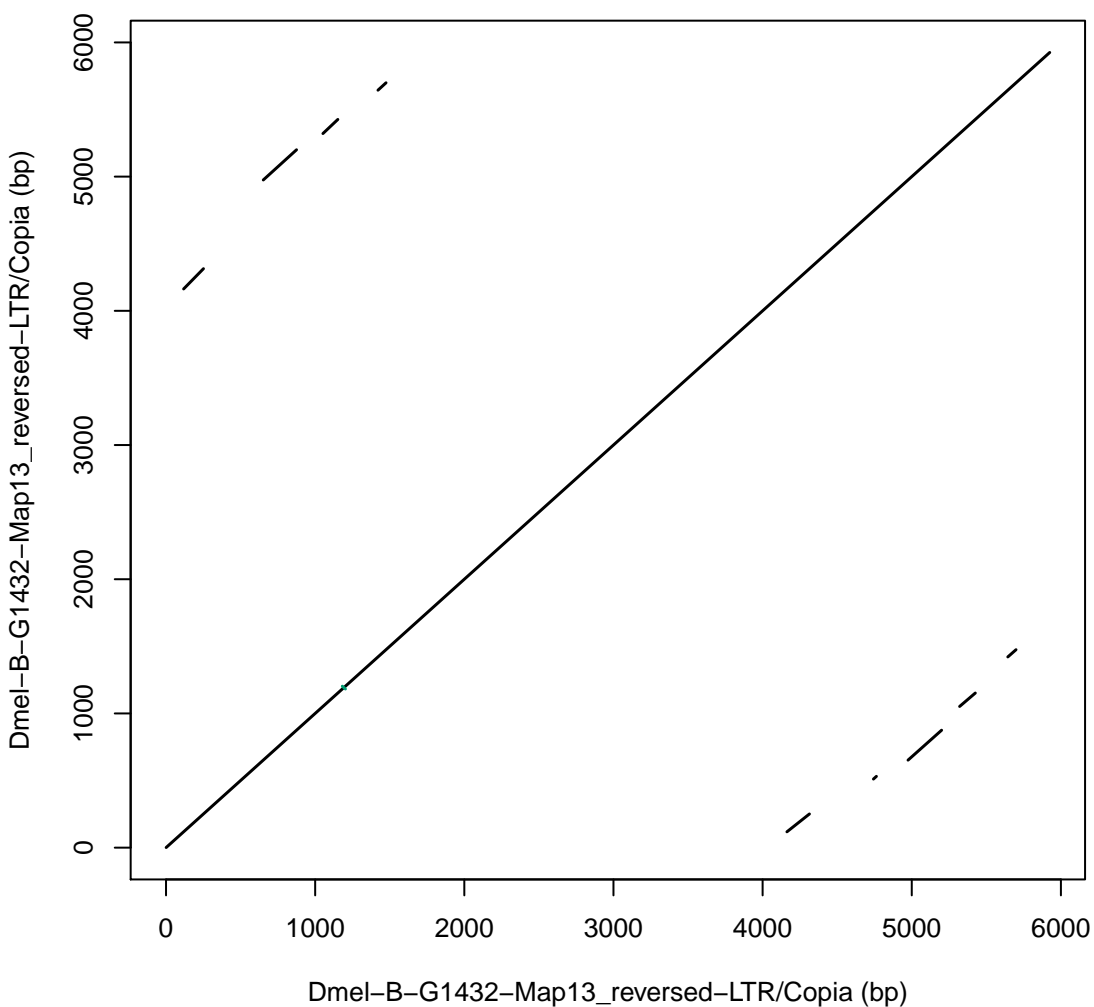
TE: Dmel-B-G1432-Map13_reversed-LTR/Copia
consensus size: 5925bp; fragments: 90; full length: 0 (>=5332.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

