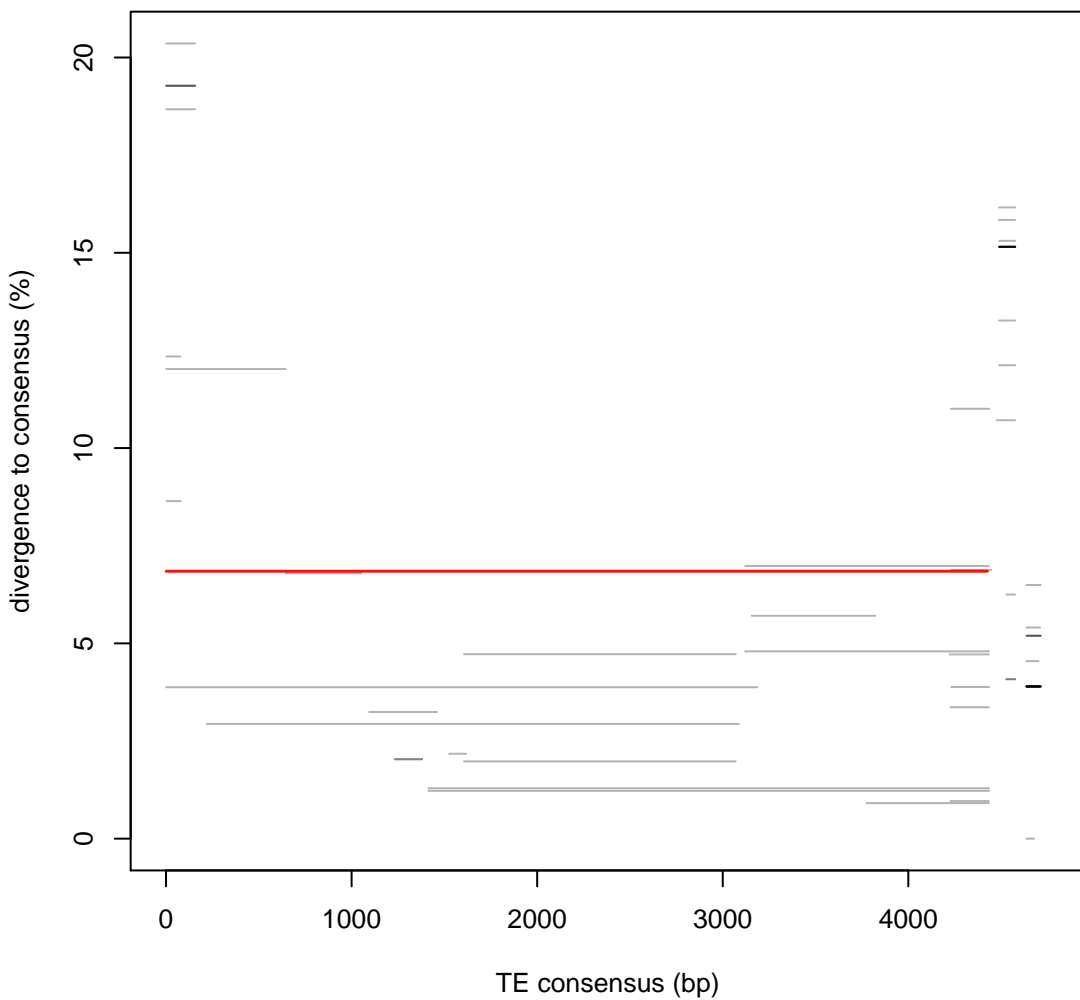
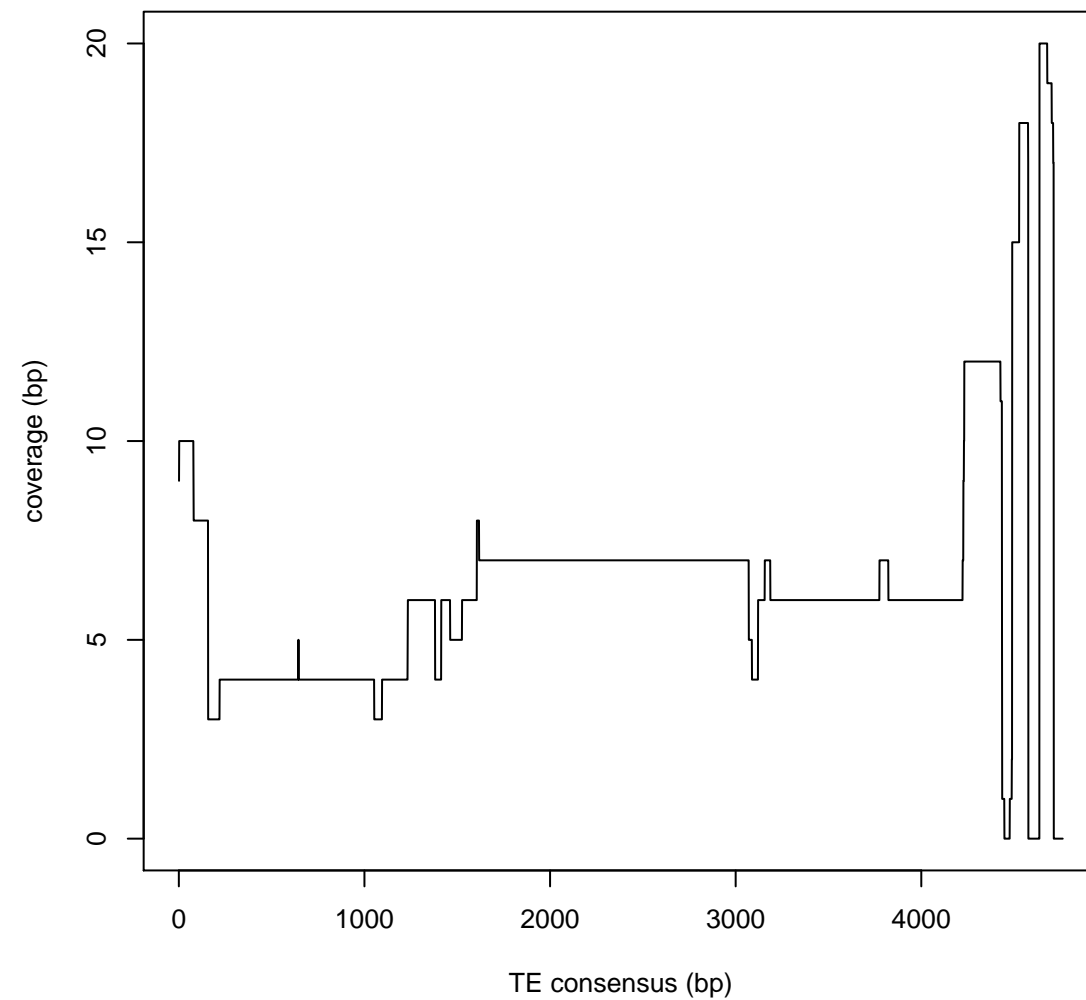


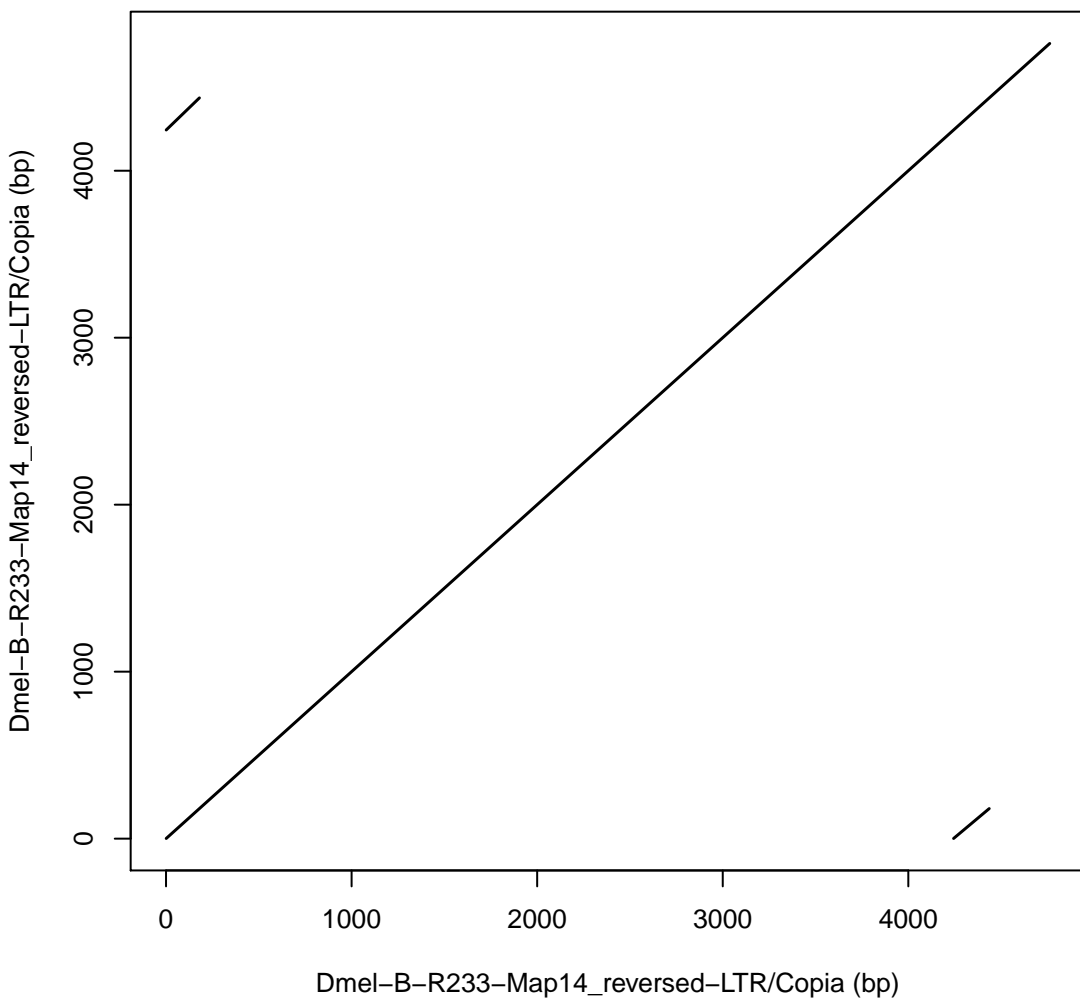
TE: Dmel-B-R233-Map14\_reversed-LTR/Copia  
consensus size: 4762bp; fragments: 68; full length: 1 (>=4285.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

