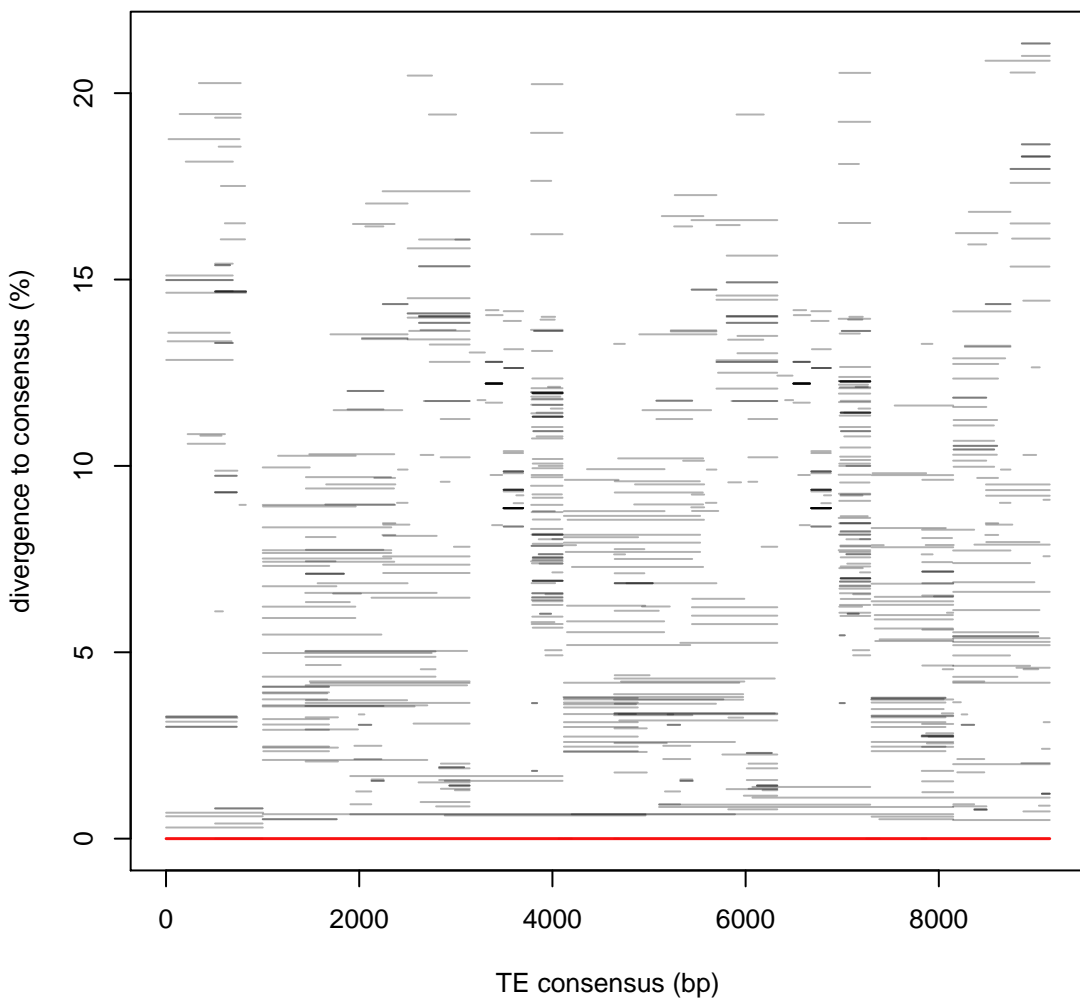
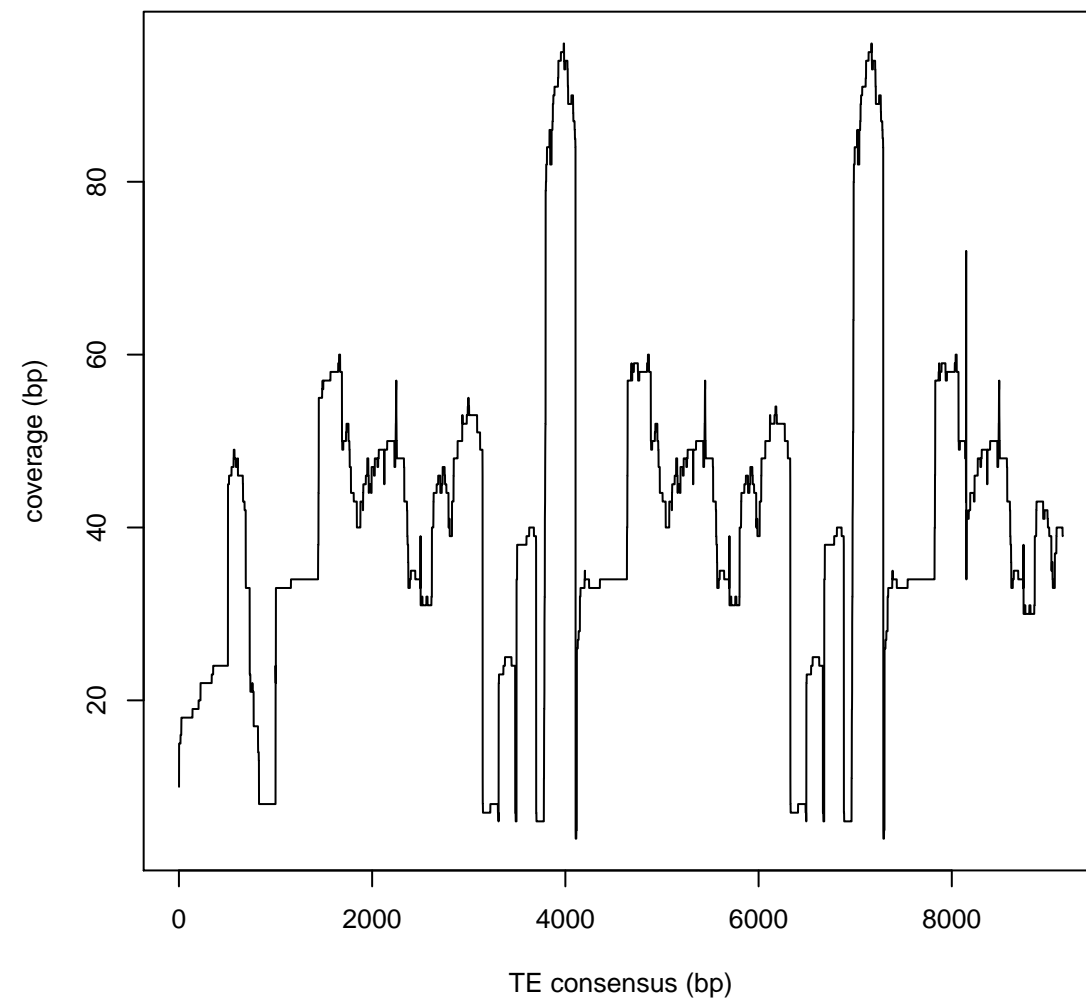


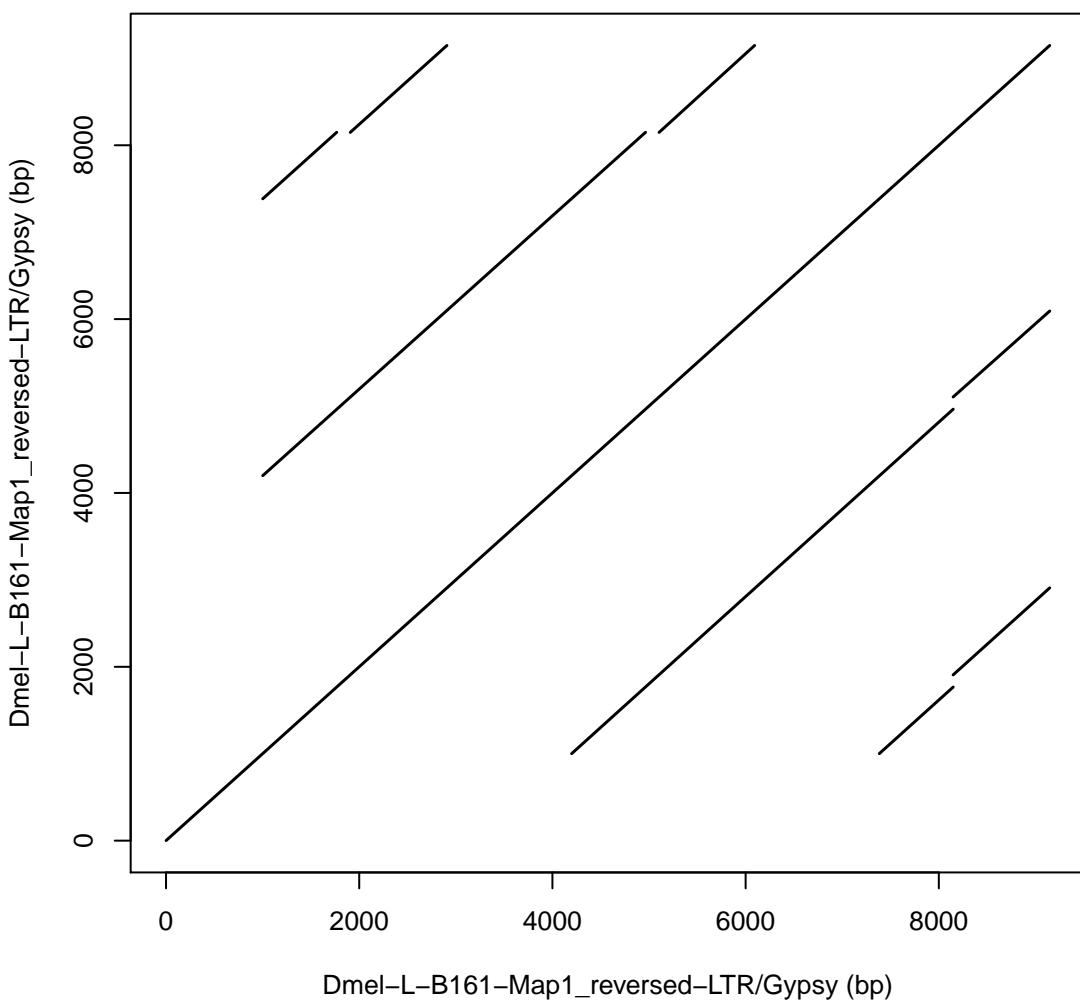
TE: Dmel-L-B161-Map1_reversed-LTR/Gypsy
consensus size: 9148bp; fragments: 798; full length: 1 (>=8233.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

