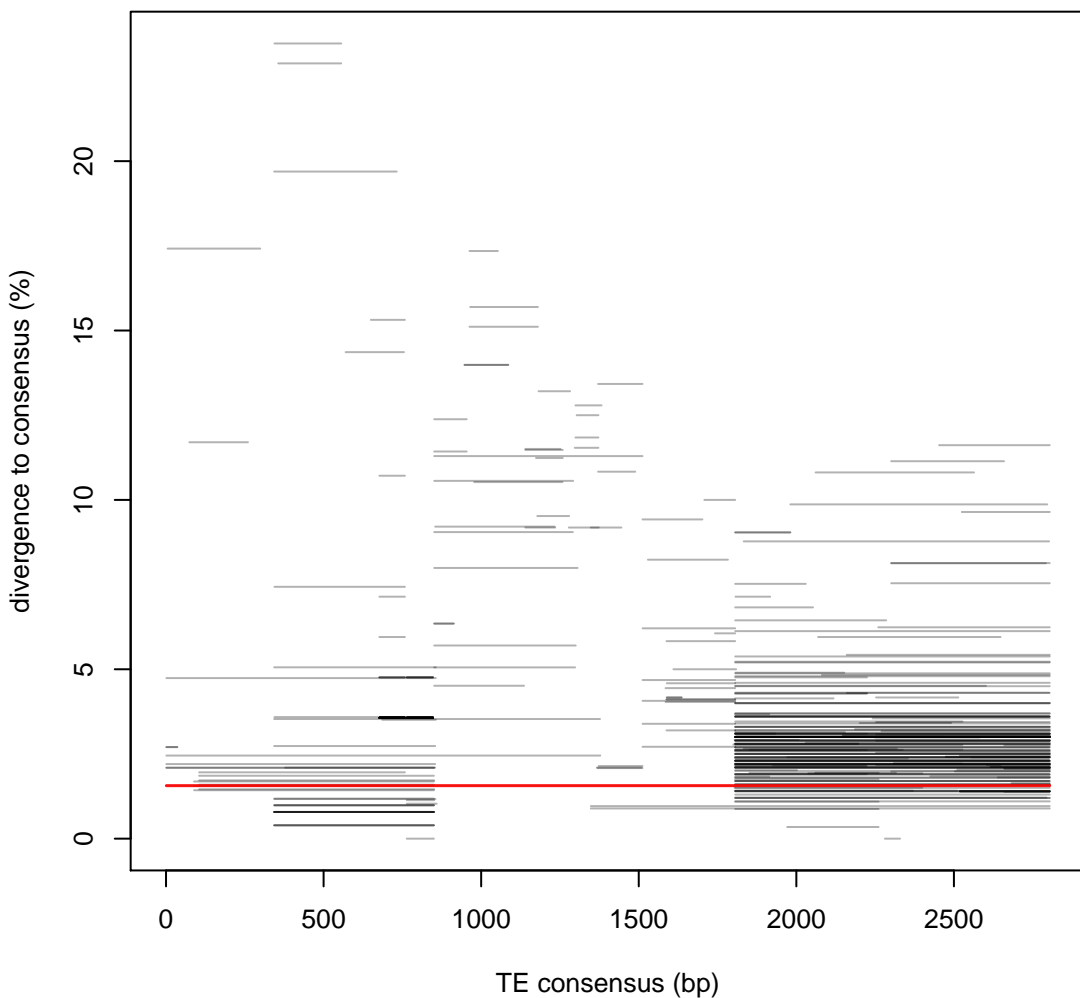
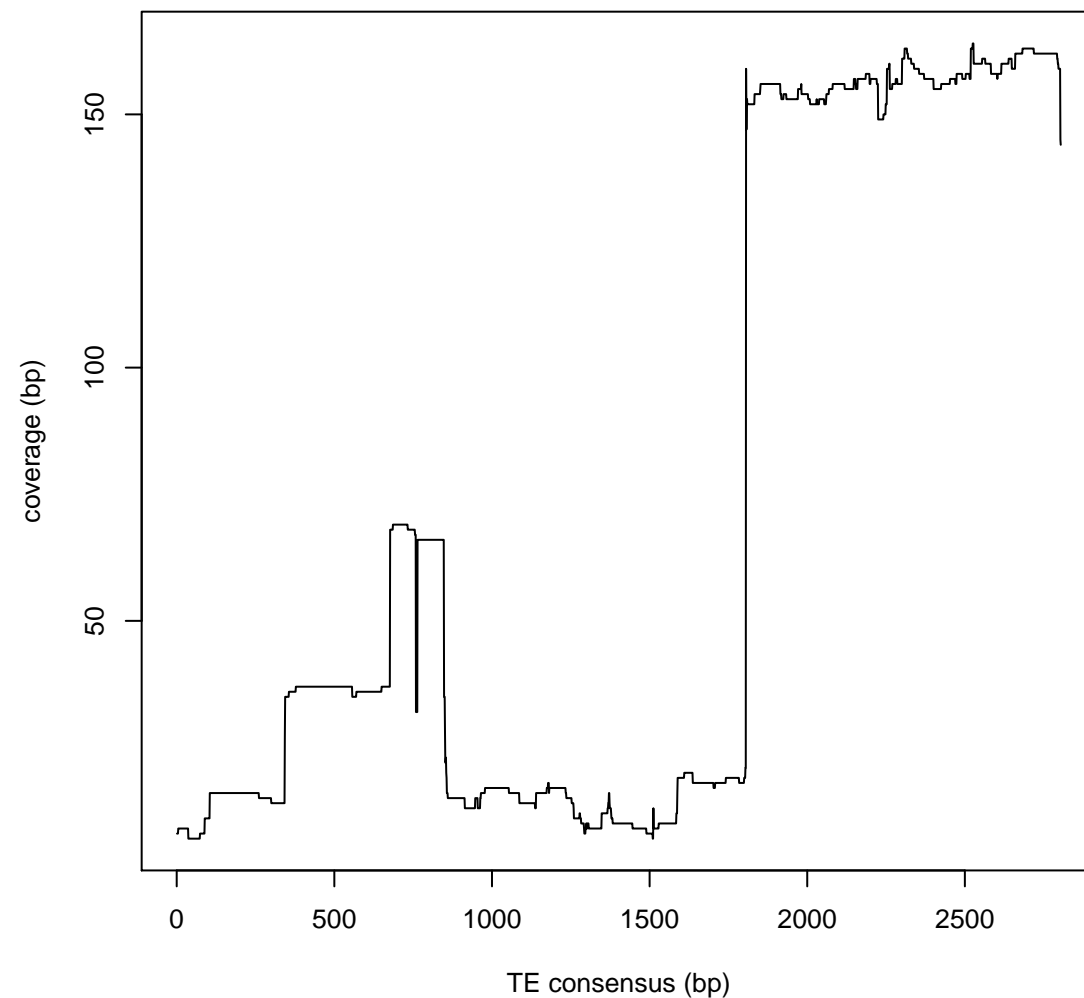


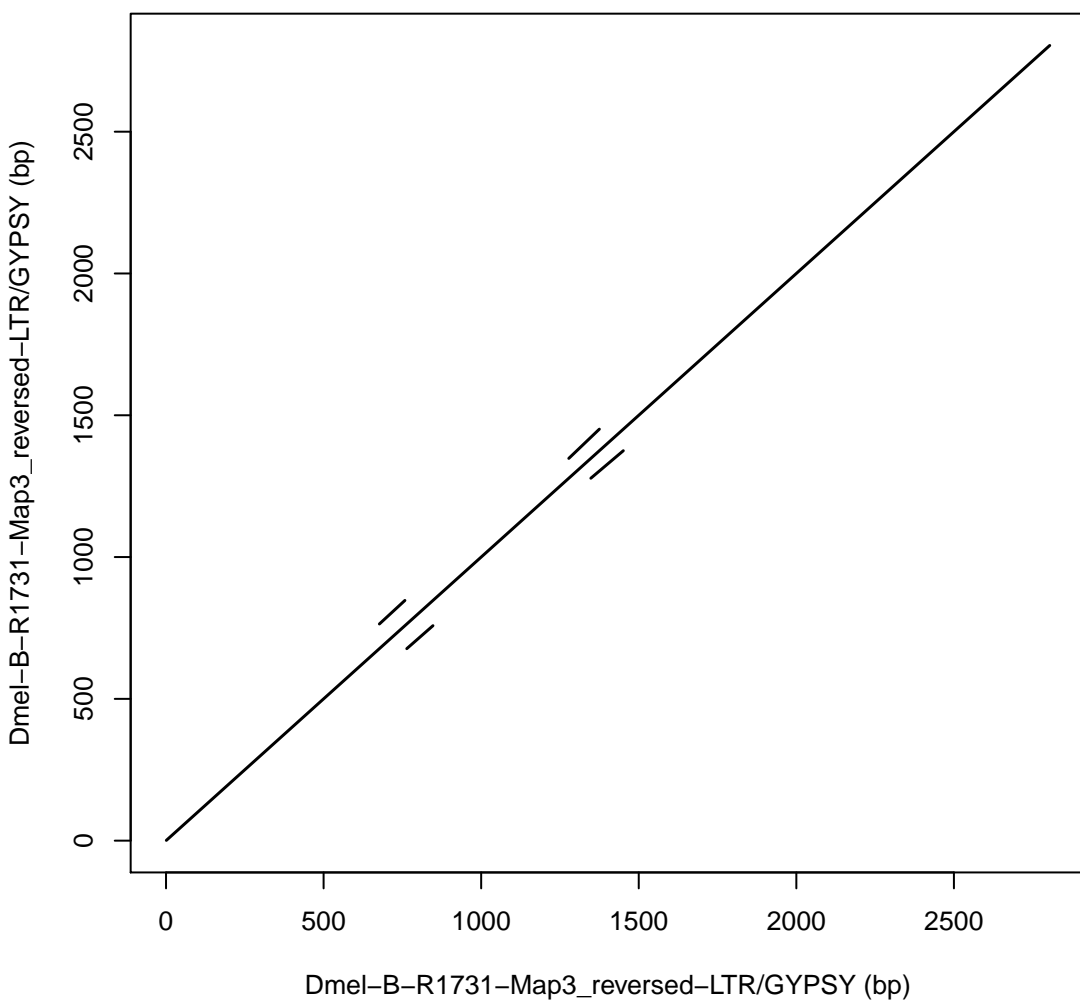
TE: Dmel-B-R1731-Map3_reversed-LTR/GYPHY
consensus size: 2804bp; fragments: 377; full length: 1 (>=2523.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

