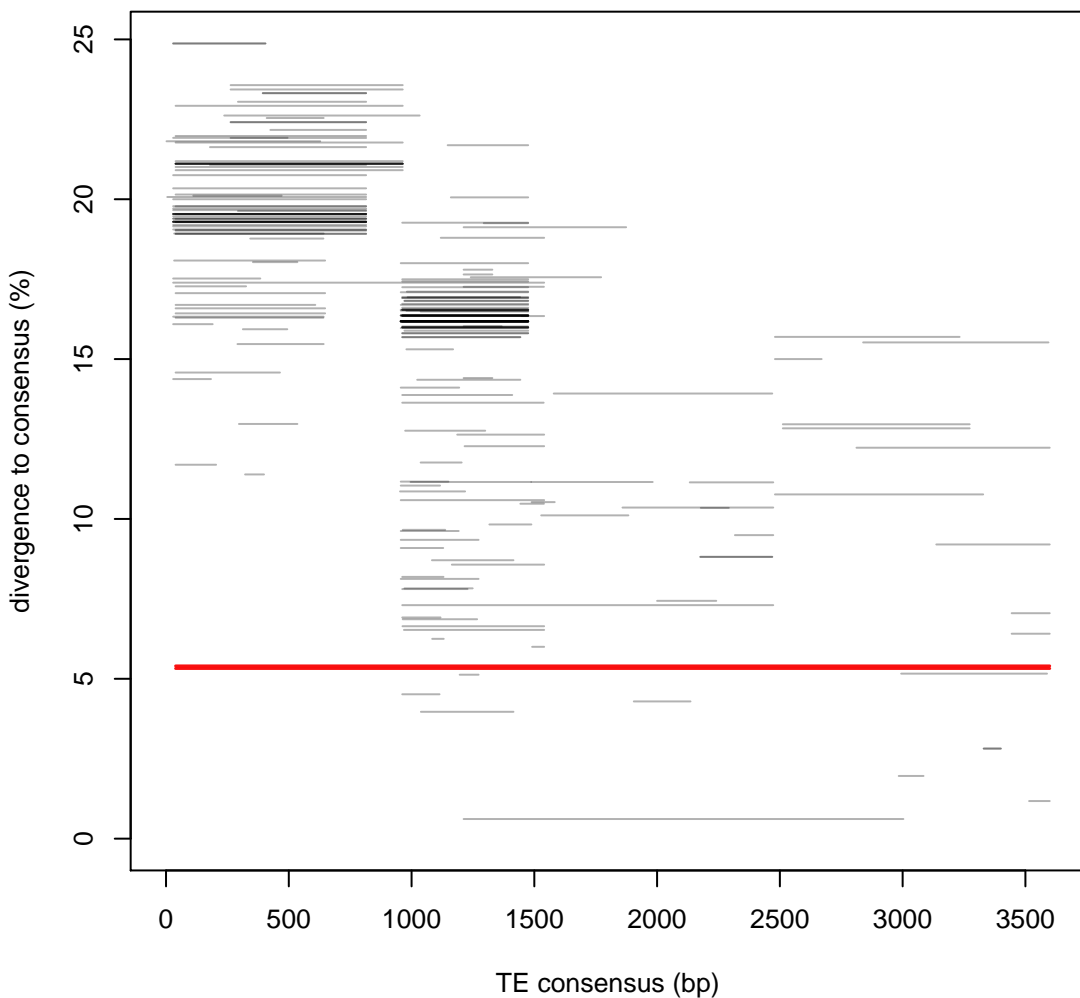
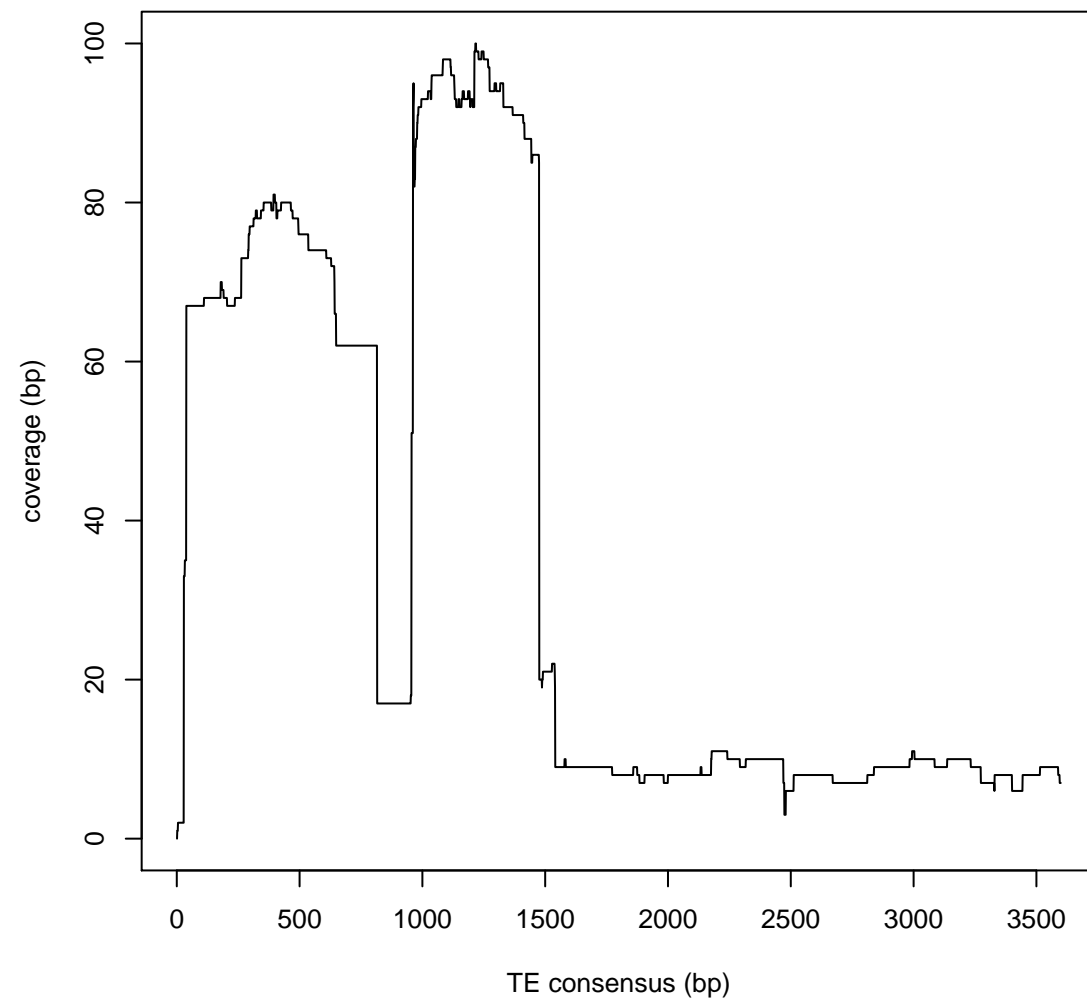


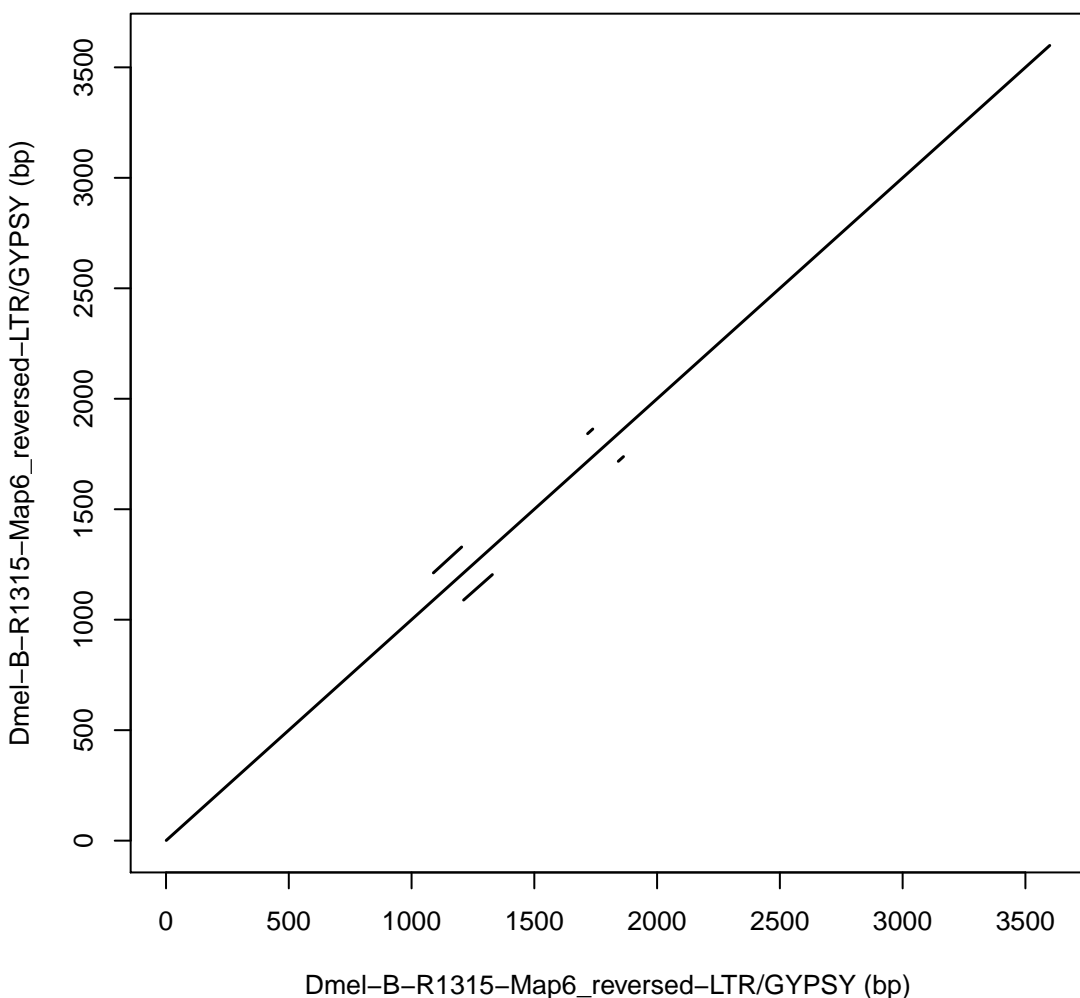
TE: Dmel-B-R1315-Map6_reversed-LTR/GYP
consensus size: 3599bp; fragments: 230; full length: 2 (>=3239.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

