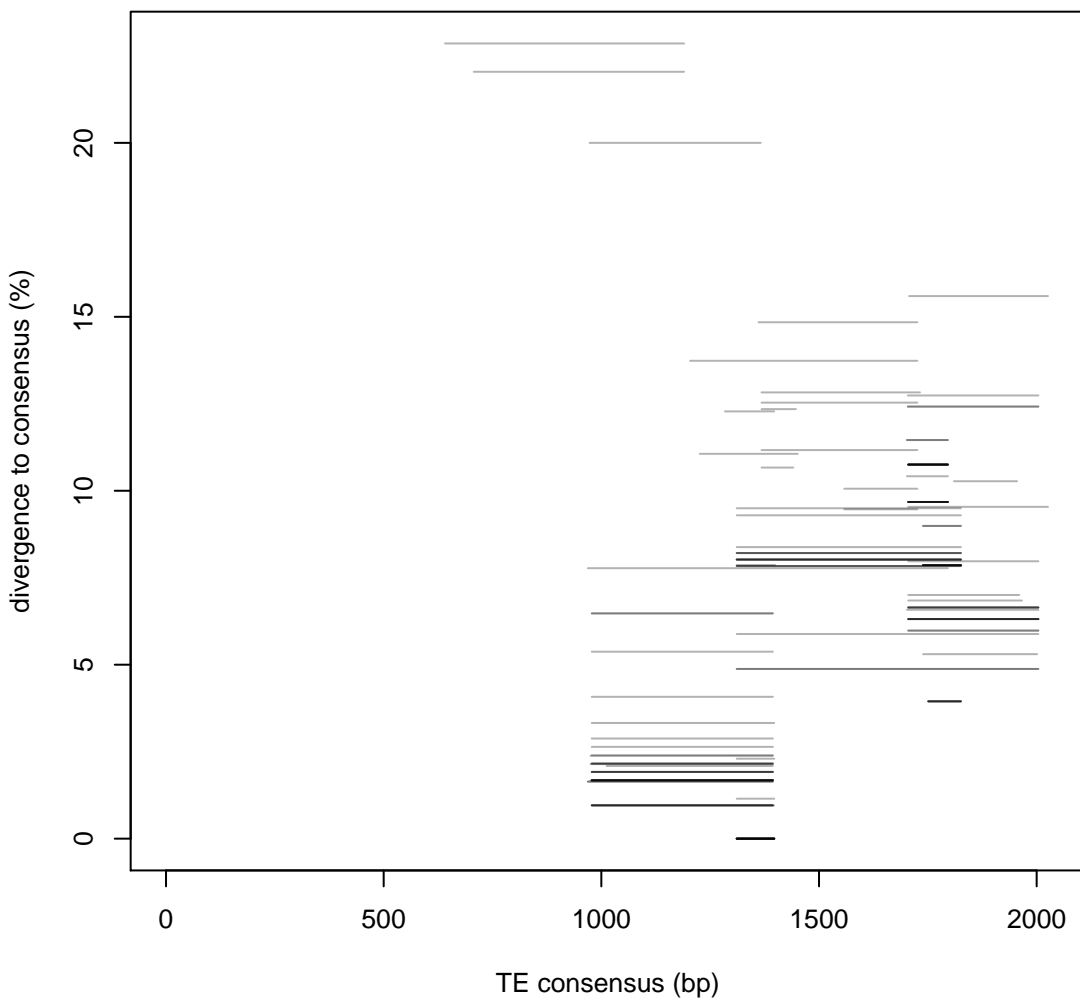
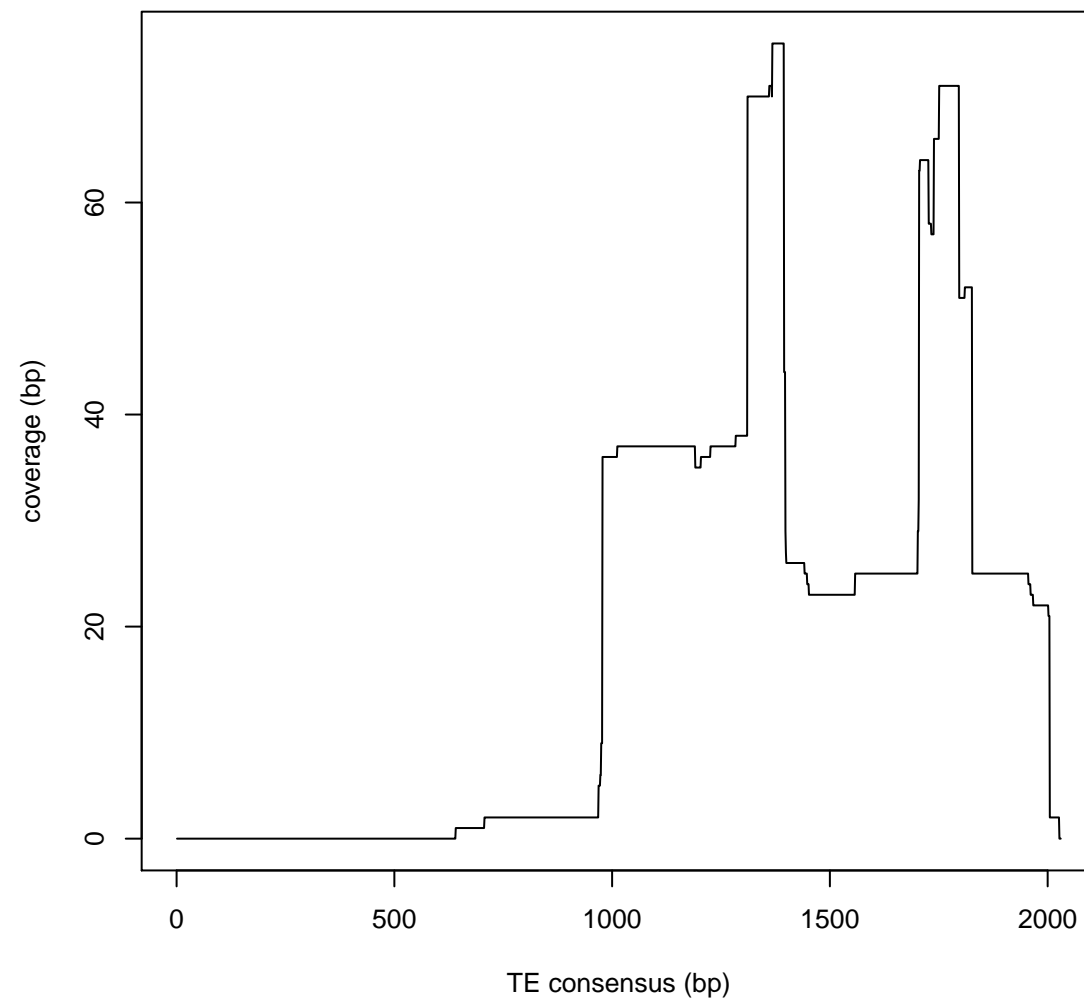


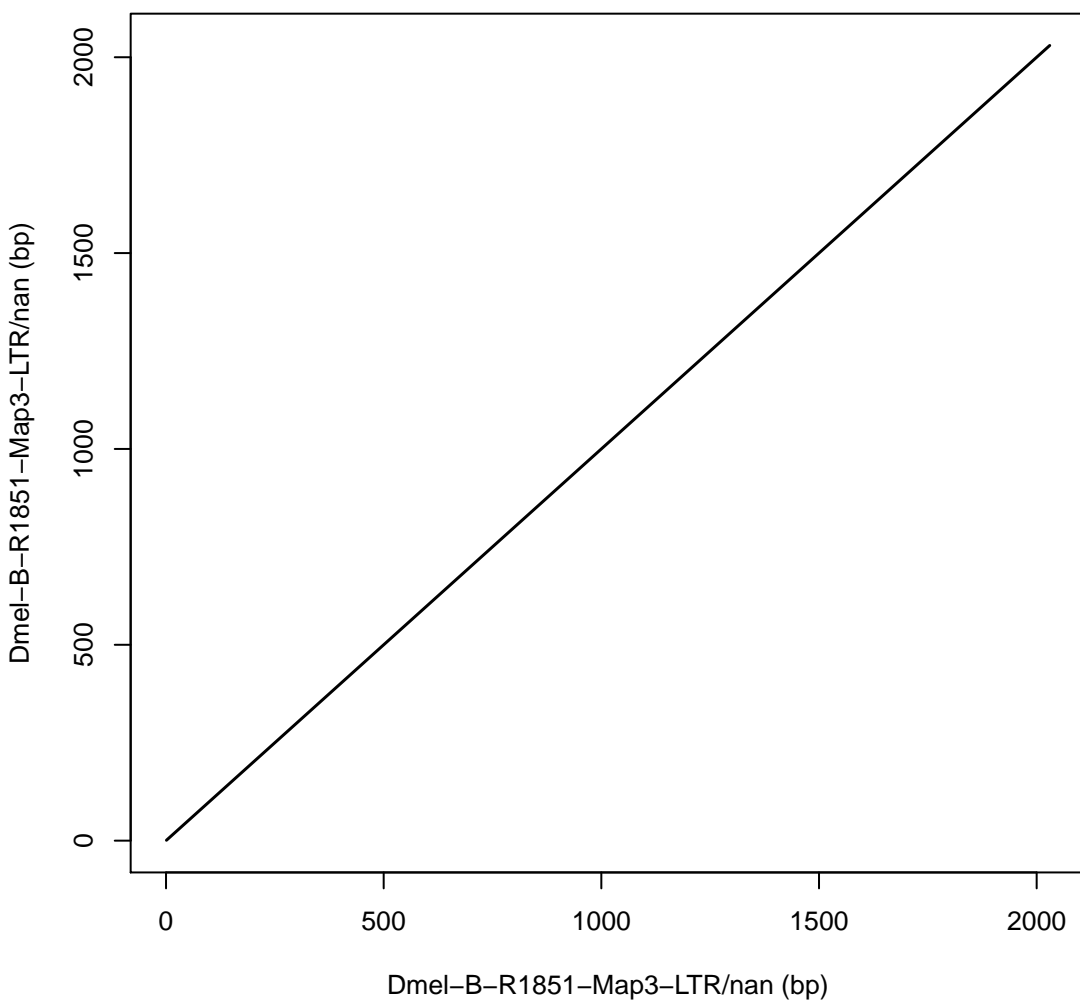
TE: Dmel-B-R1851-Map3-LTR/nan
consensus size: 2030bp; fragments: 134; full length: 0 (>=1827bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

