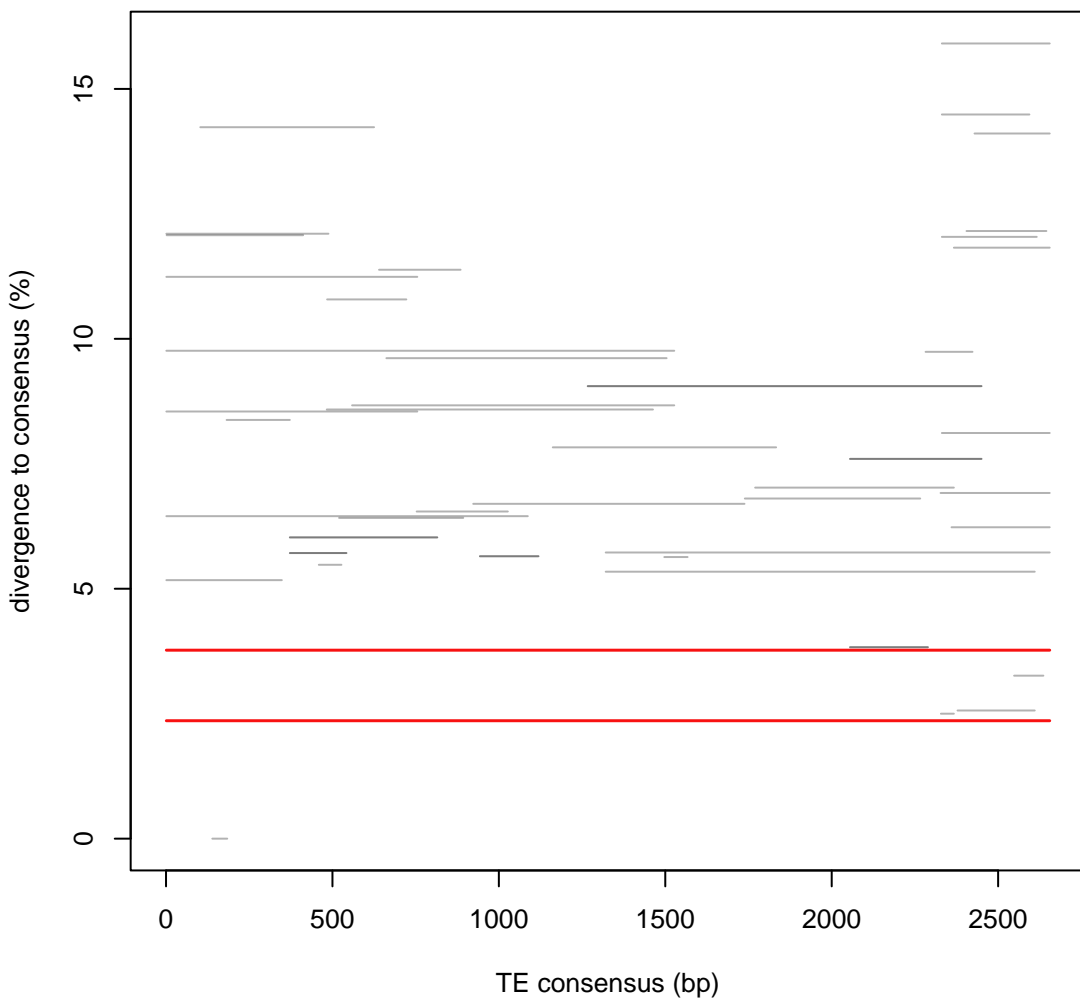
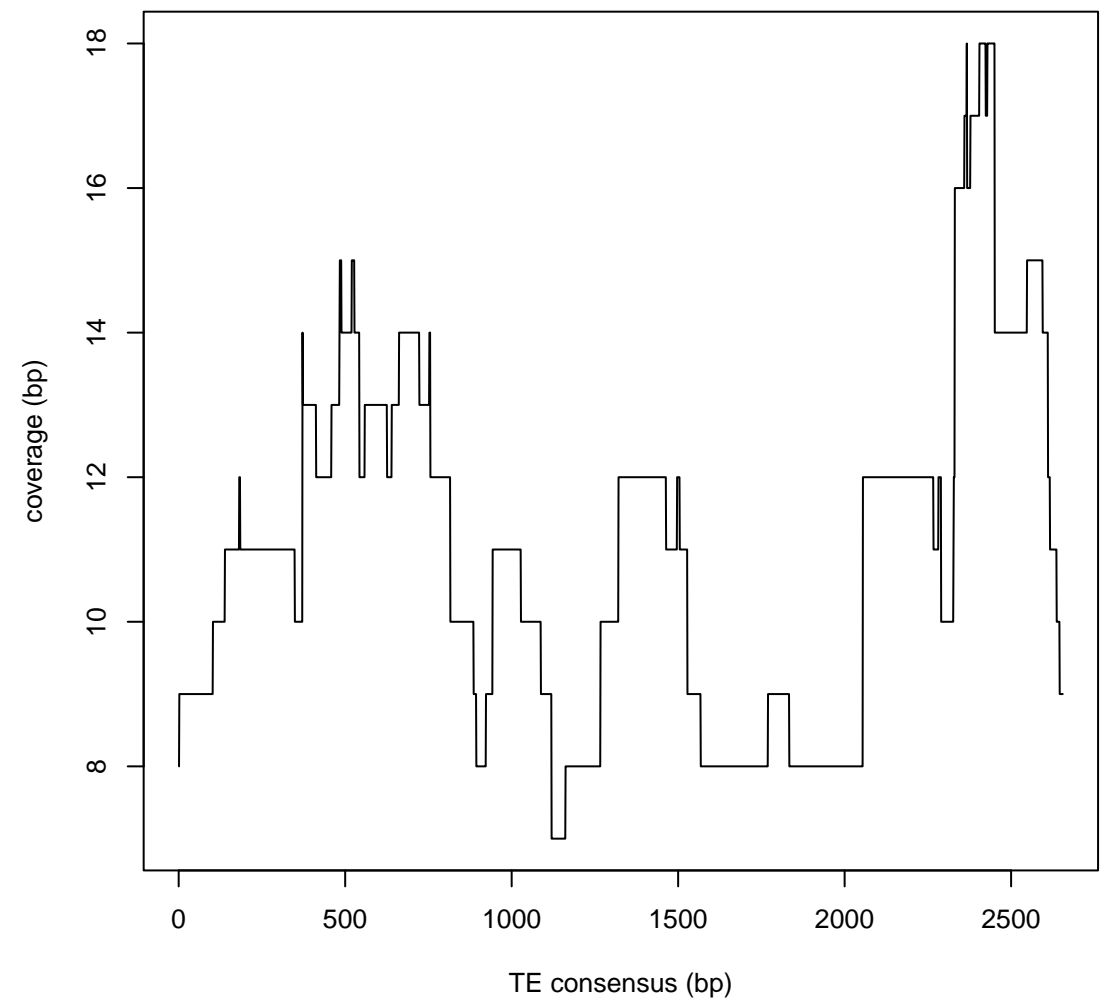


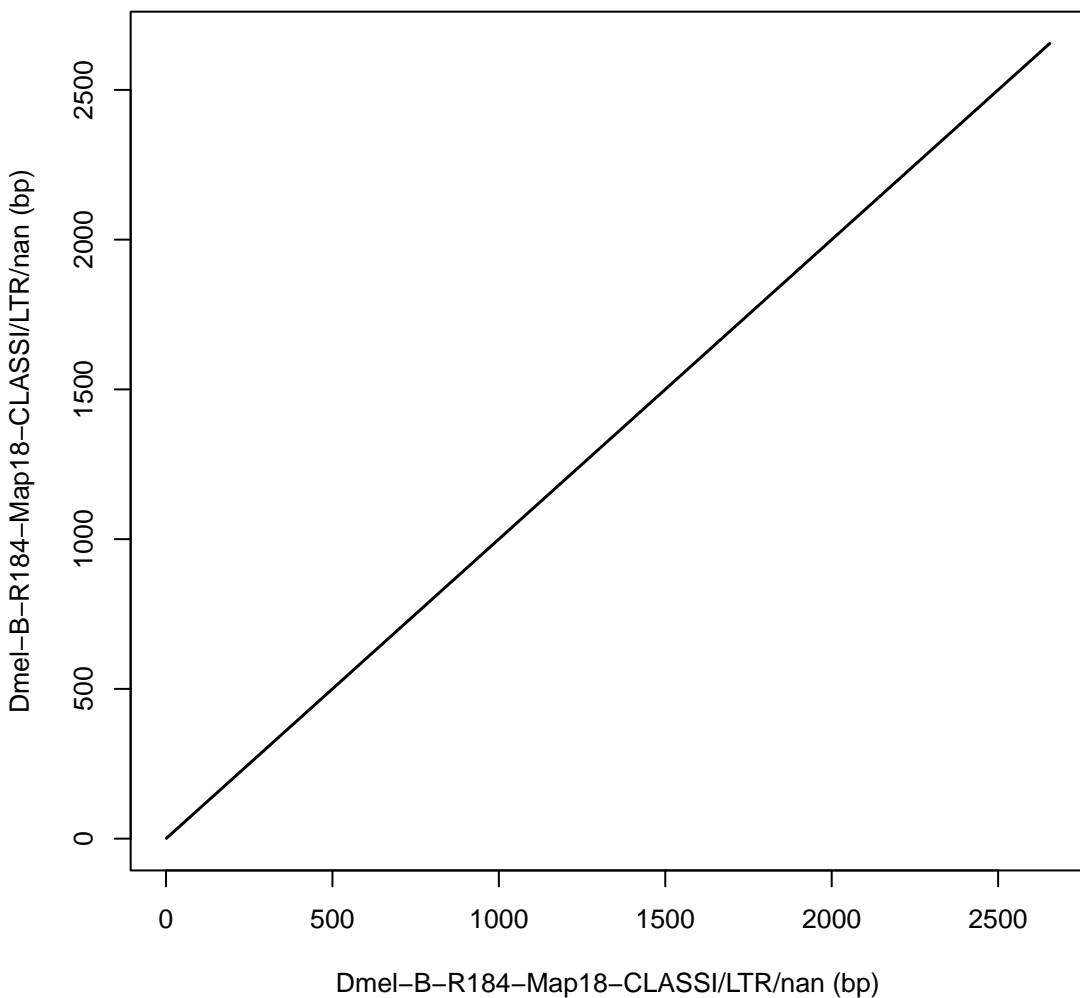
TE: Dmel-B-R184-Map18-CLASSI/LTR/nan
consensus size: 2655bp; fragments: 52; full length: 2 (>=2389.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

