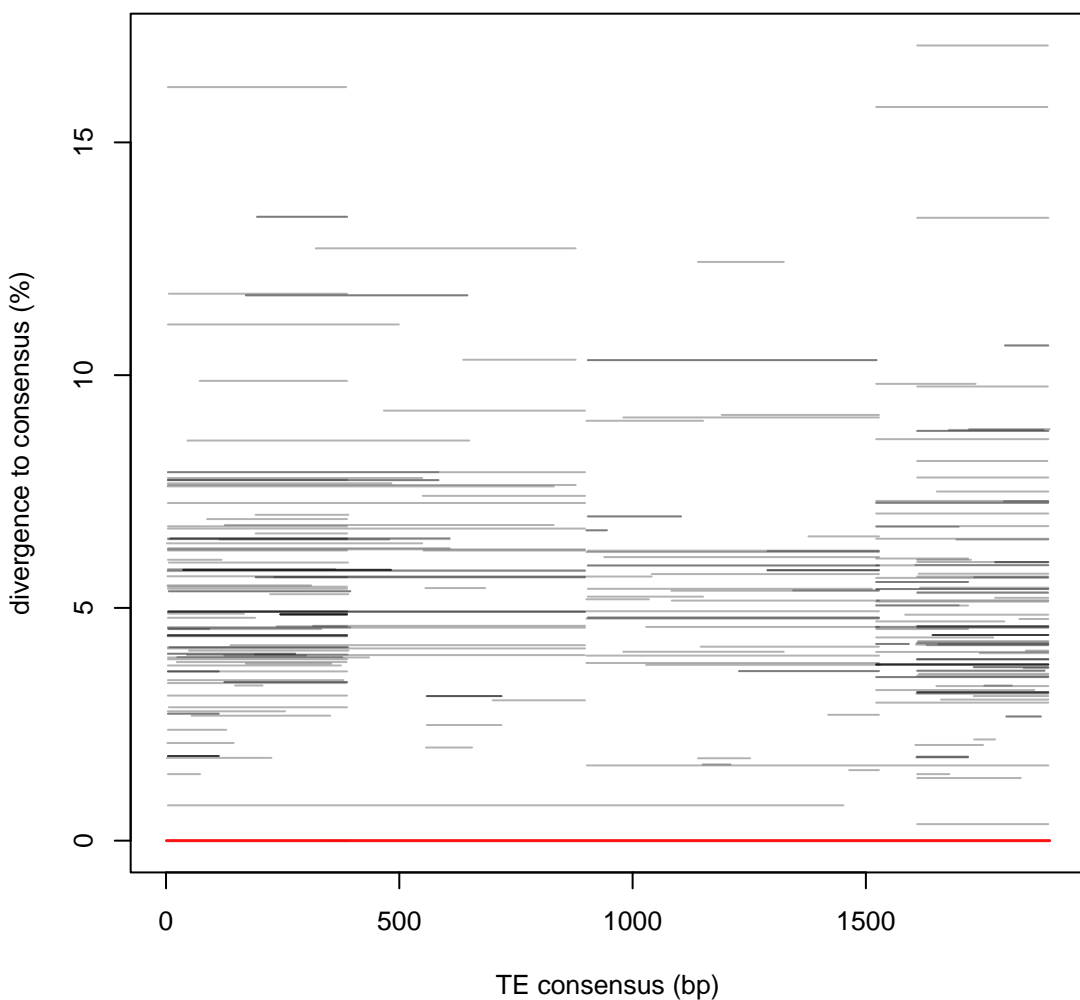
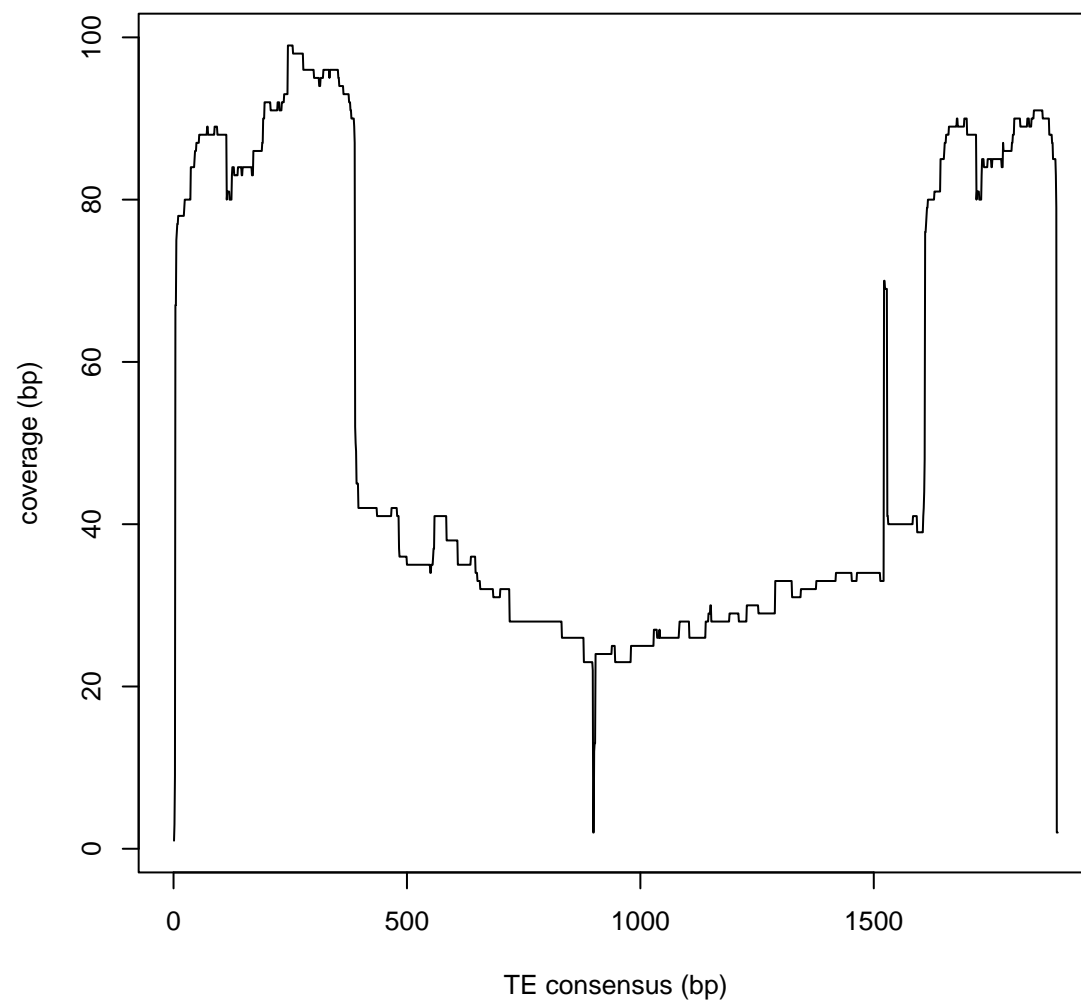


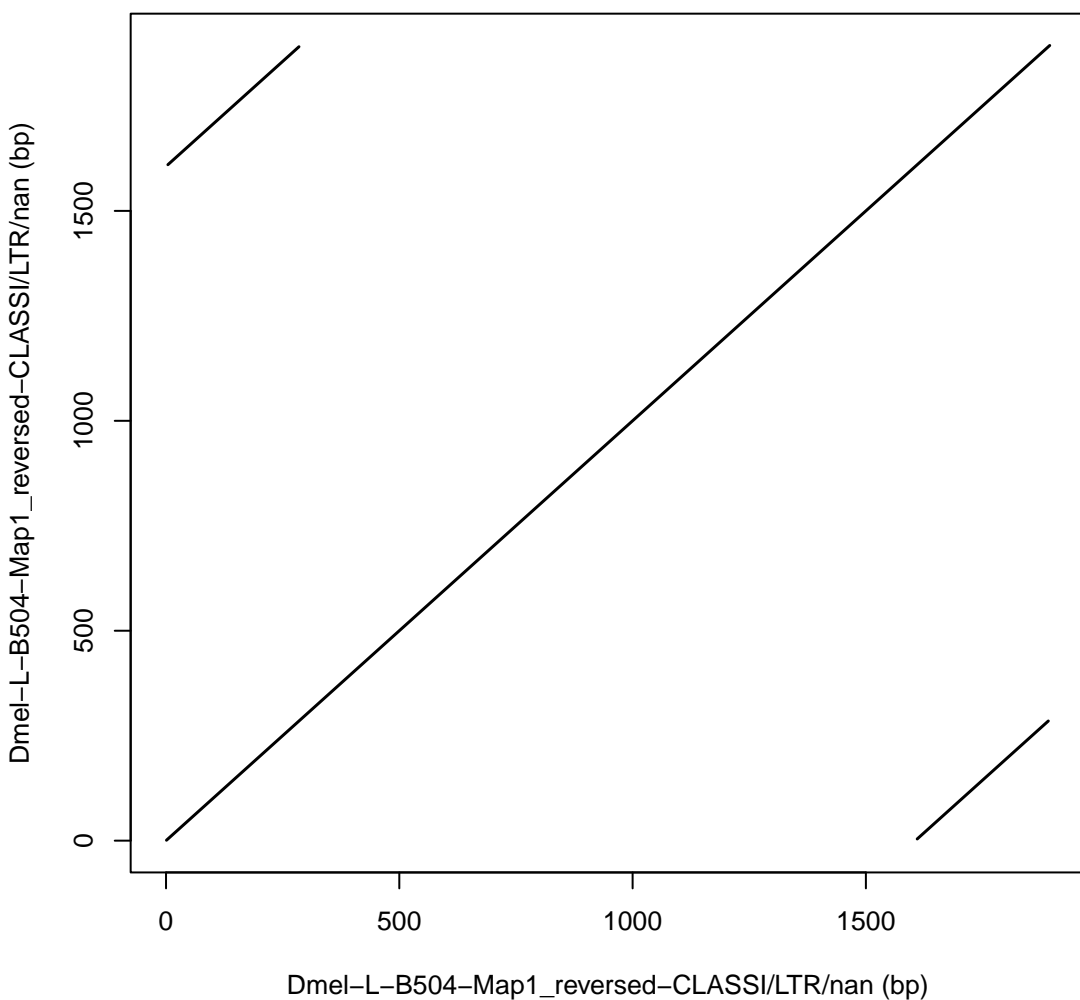
TE: Dmel-L-B504-Map1_reversed-CLASSI/LTR/nan
consensus size: 1894bp; fragments: 285; full length: 1 (>=1704.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

