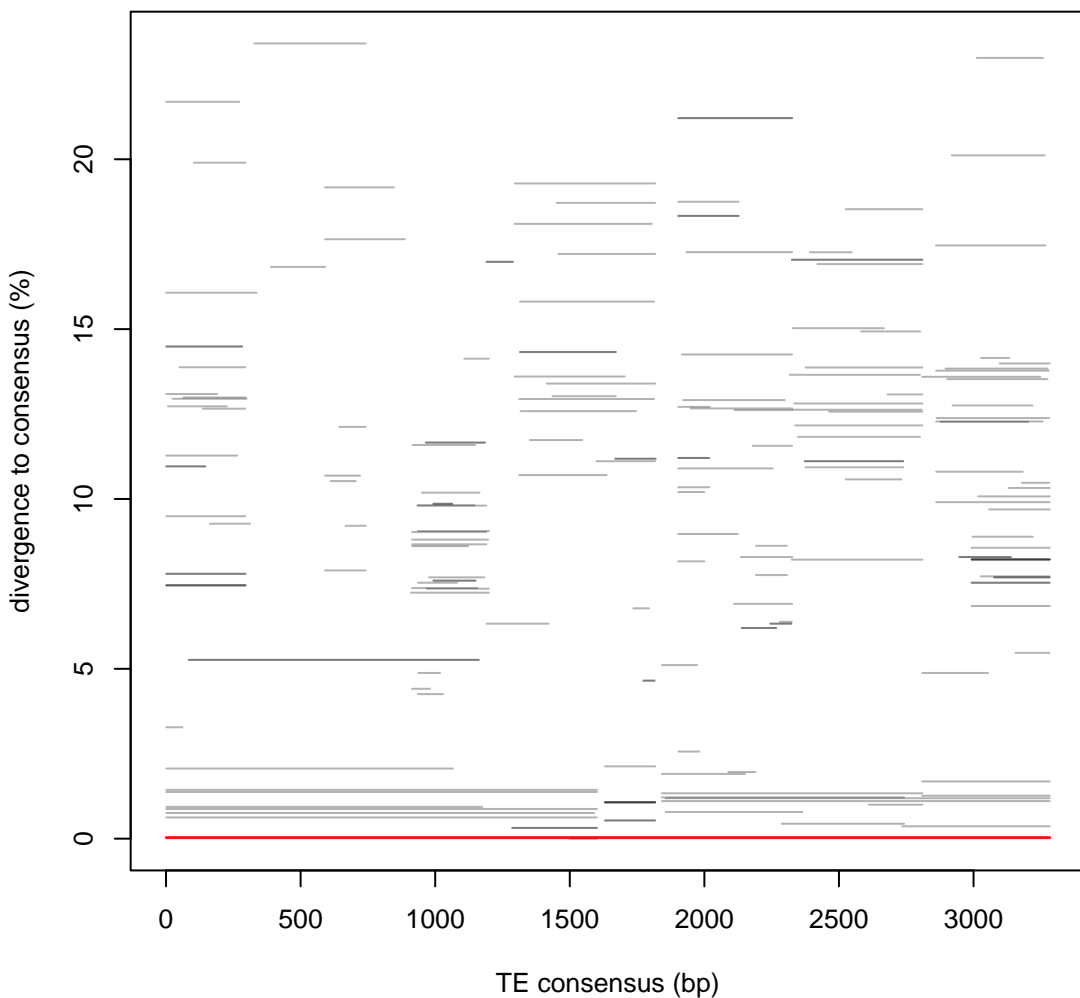
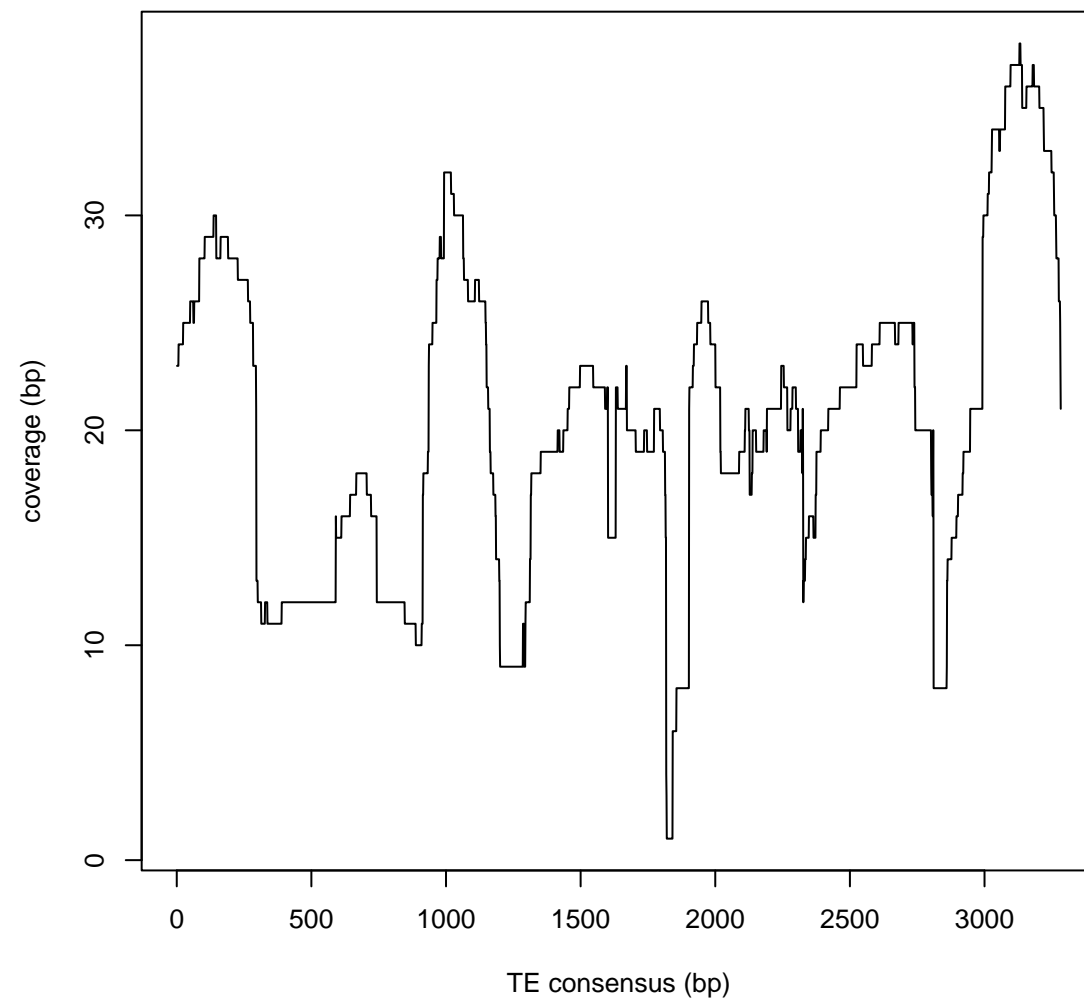


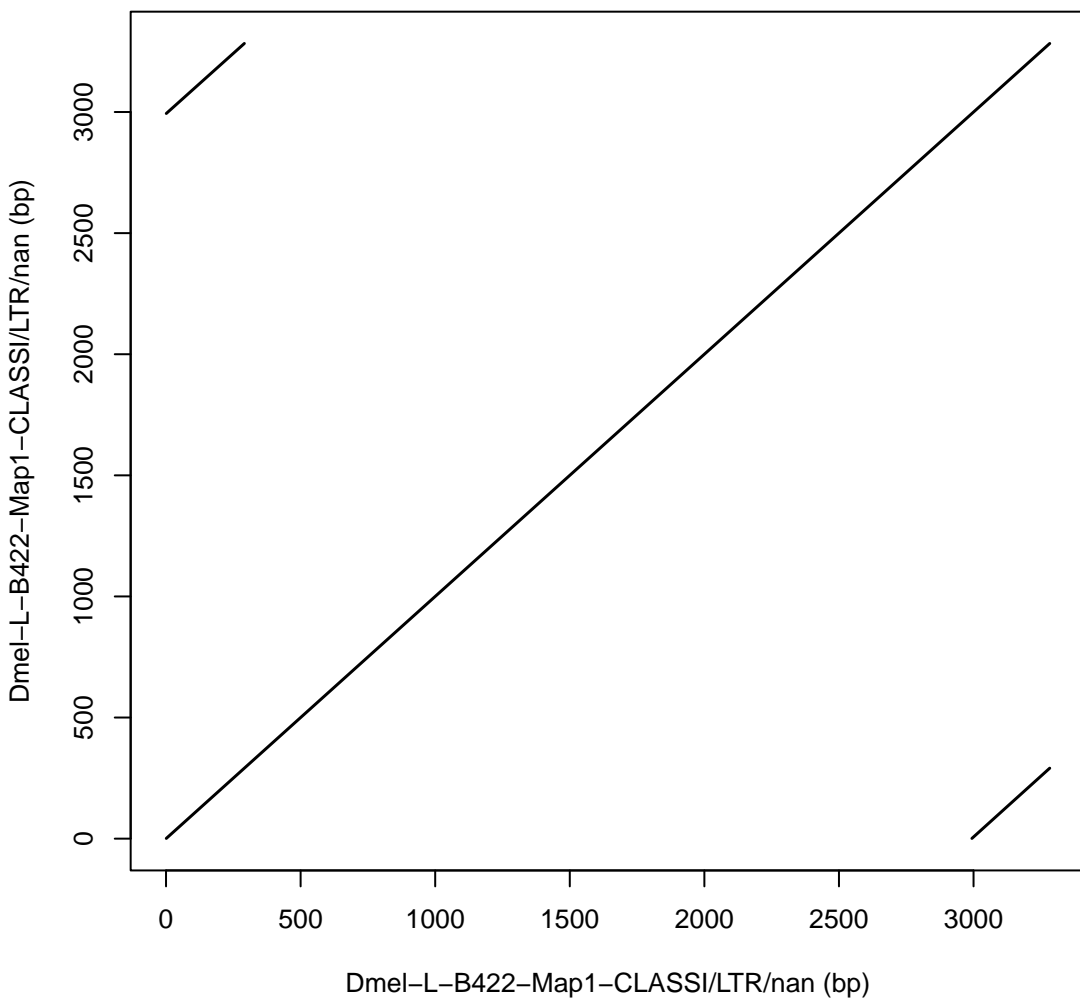
TE: Dmel-L-B422-Map1-CLASSI/LTR/nan
consensus size: 3283bp; fragments: 195; full length: 1 (>=2954.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

