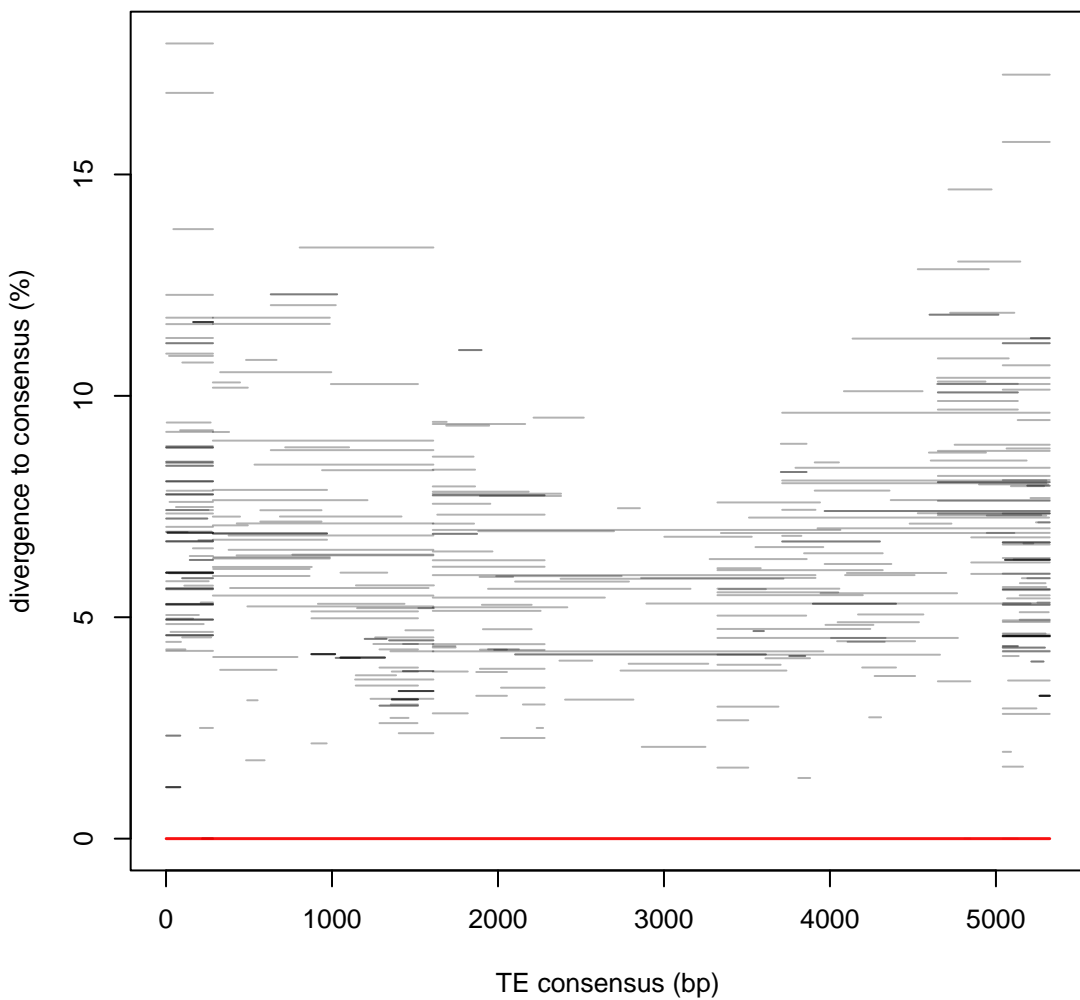
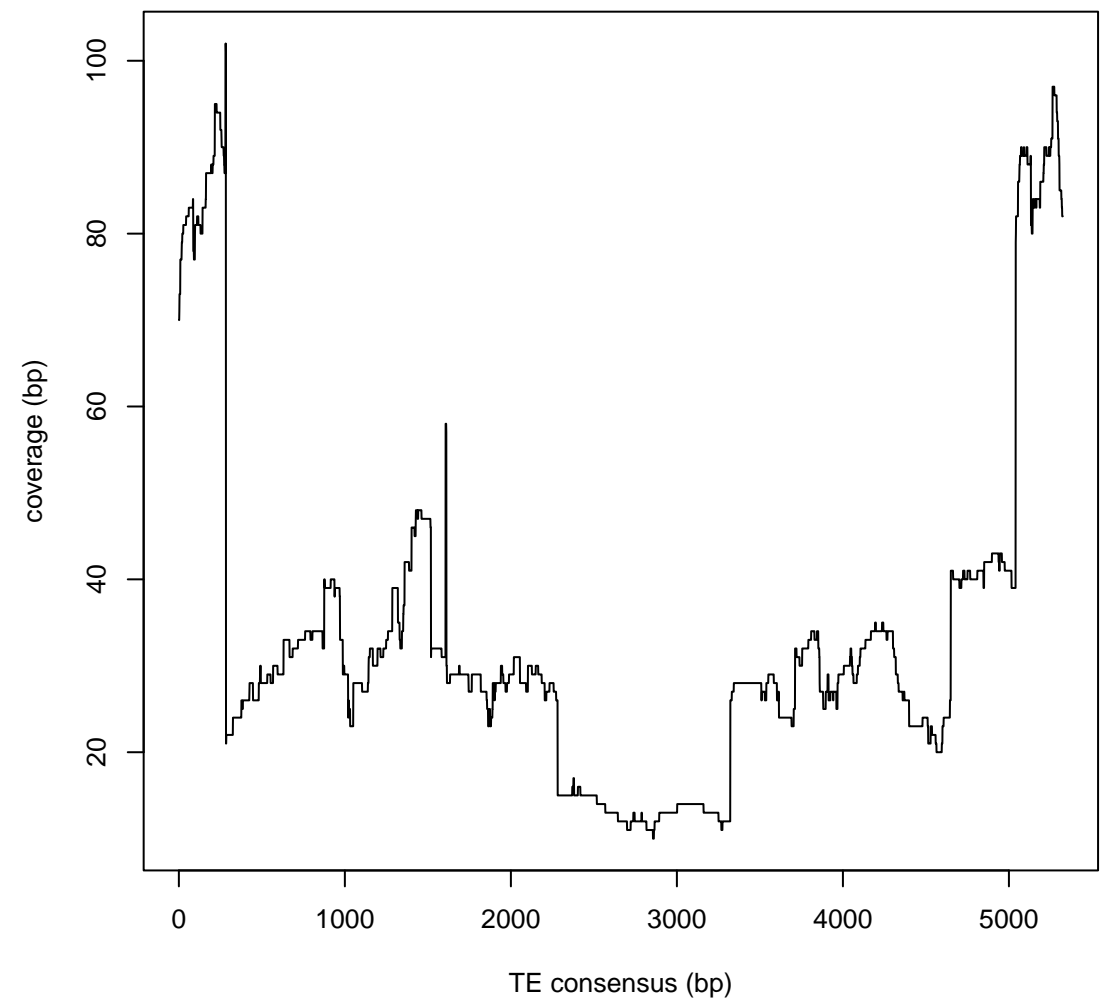


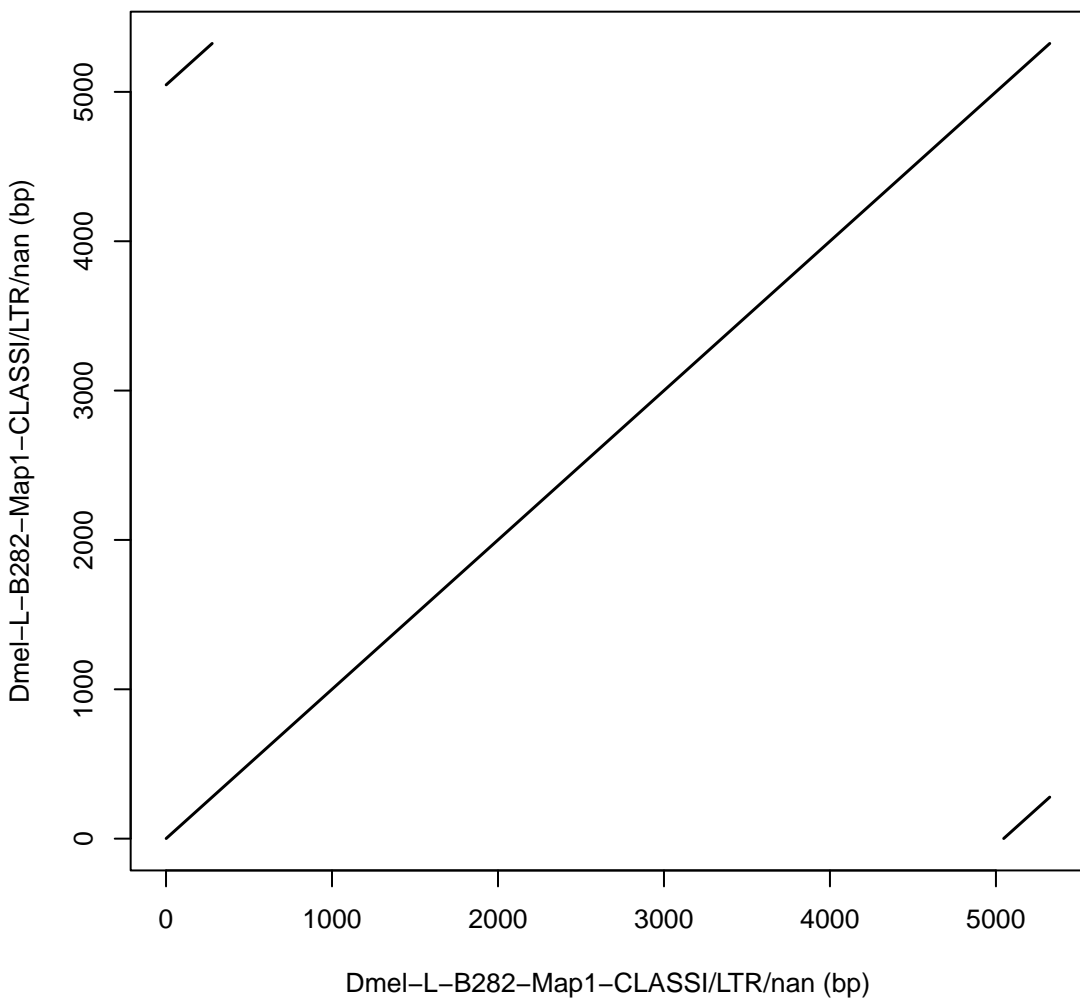
TE: Dmel-L-B282-Map1-CLASSI/LTR/nan
consensus size: 5324bp; fragments: 436; full length: 1 (>=4791.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

