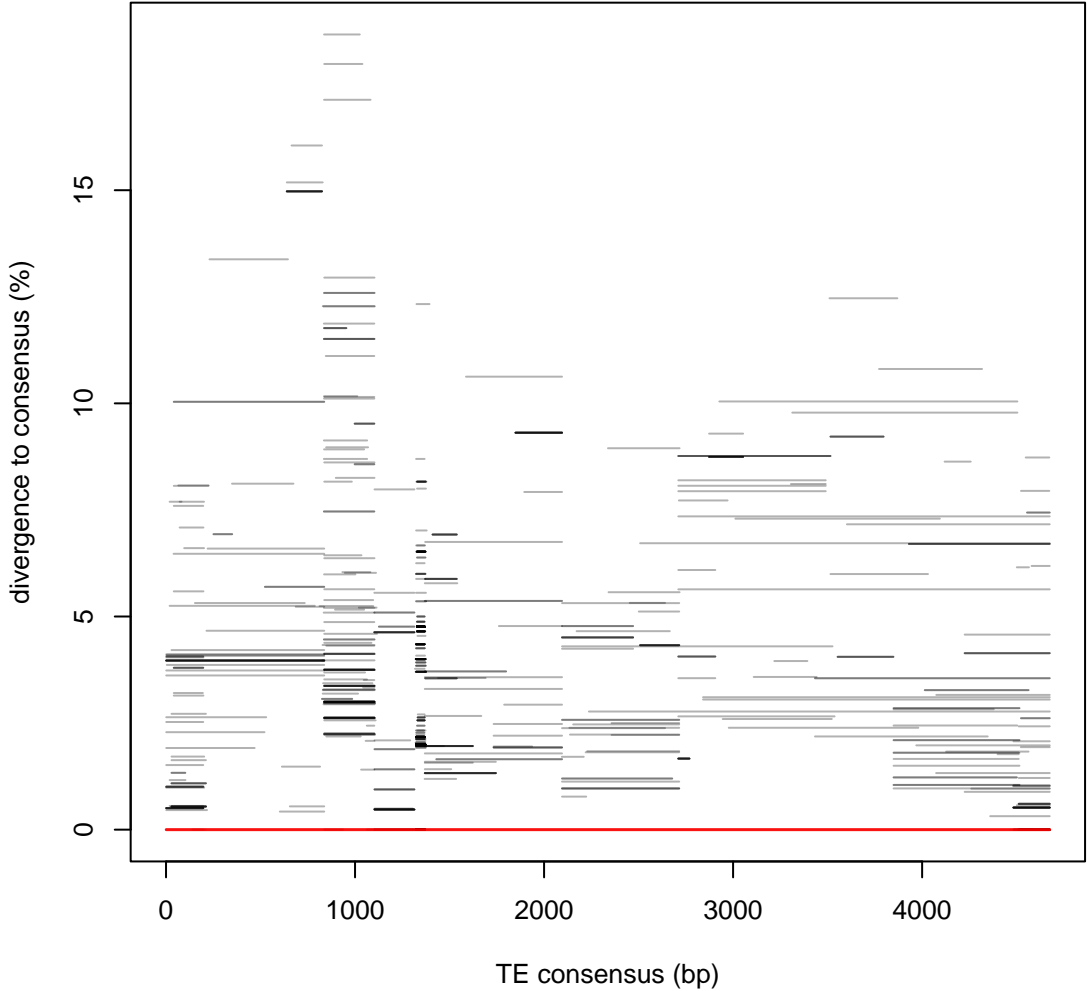
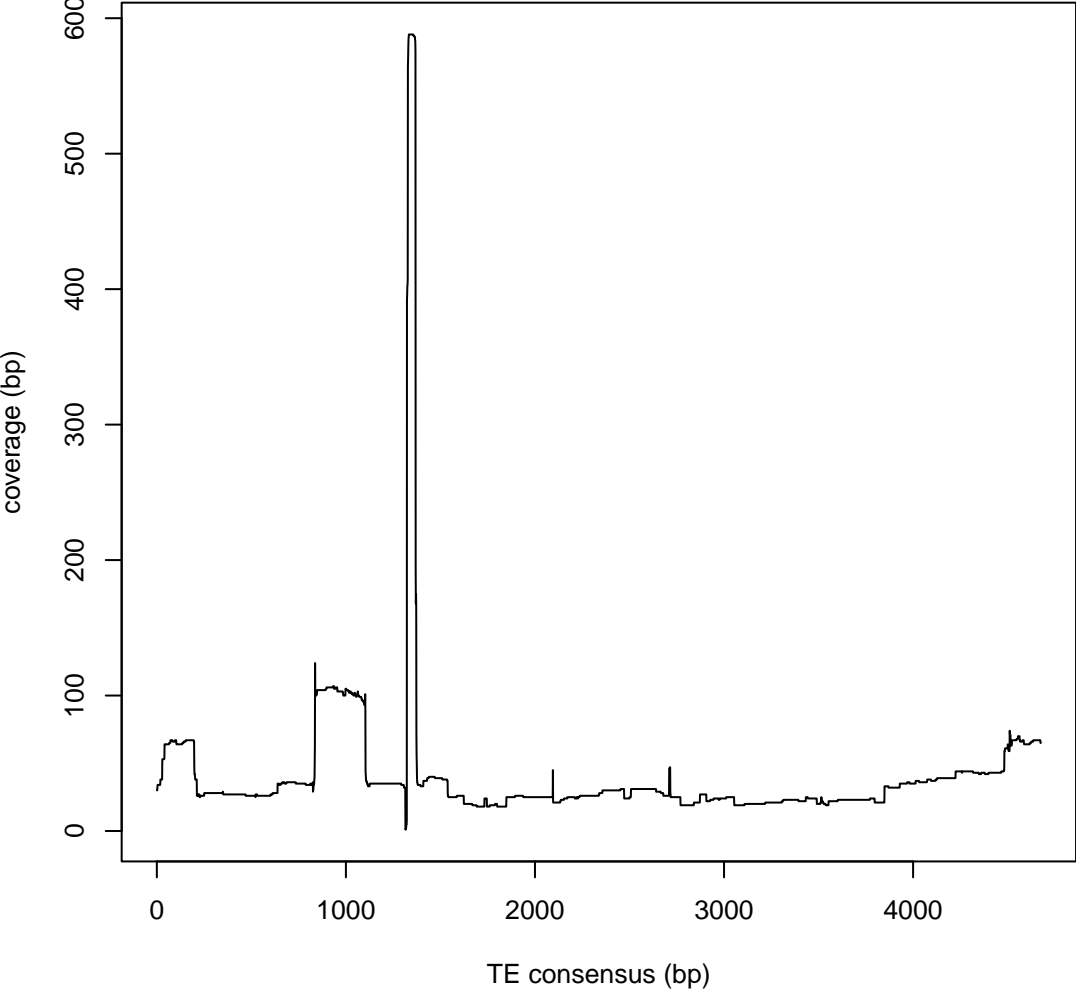


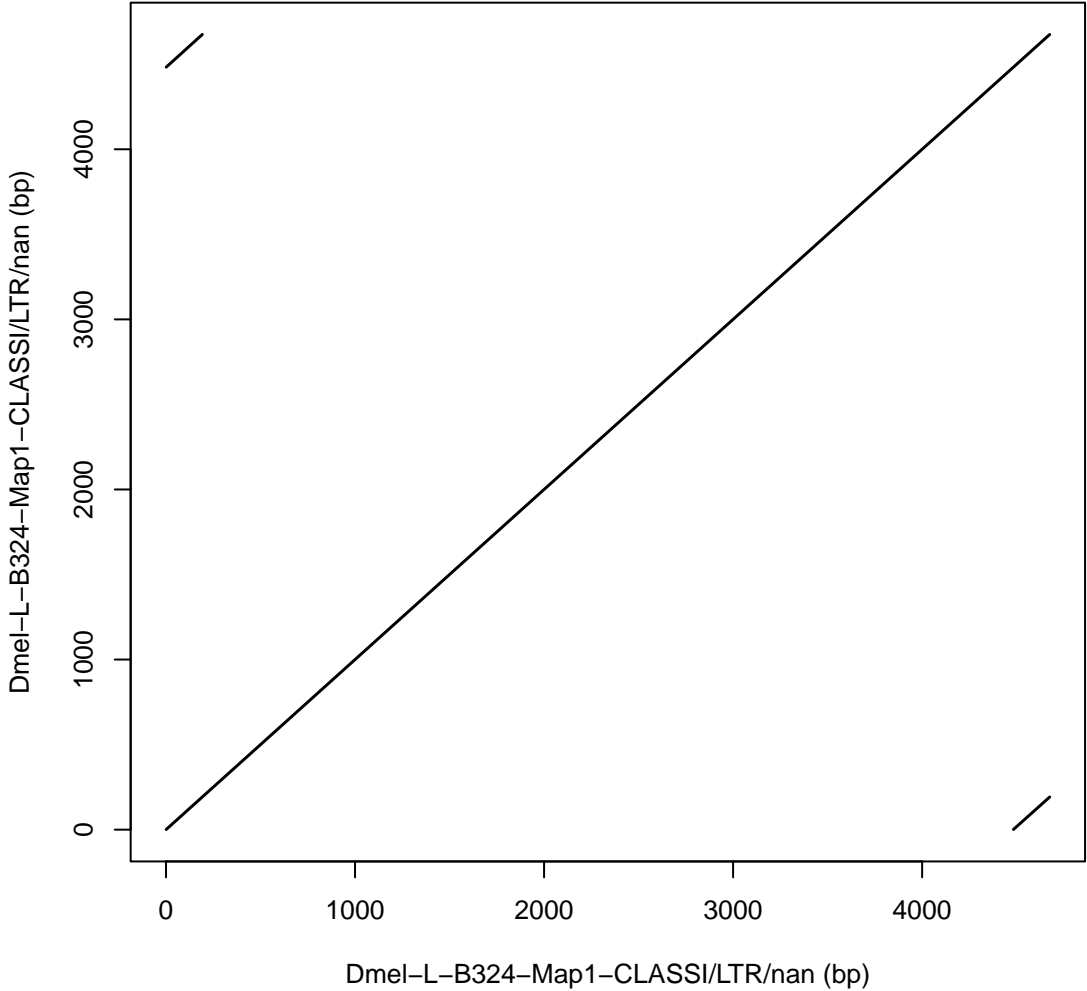
TE: Dmel-L-B324-Map1-CLASSI/LTR/nan
consensus size: 4675bp; fragments: 1057; full length: 1 (>=4207.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

