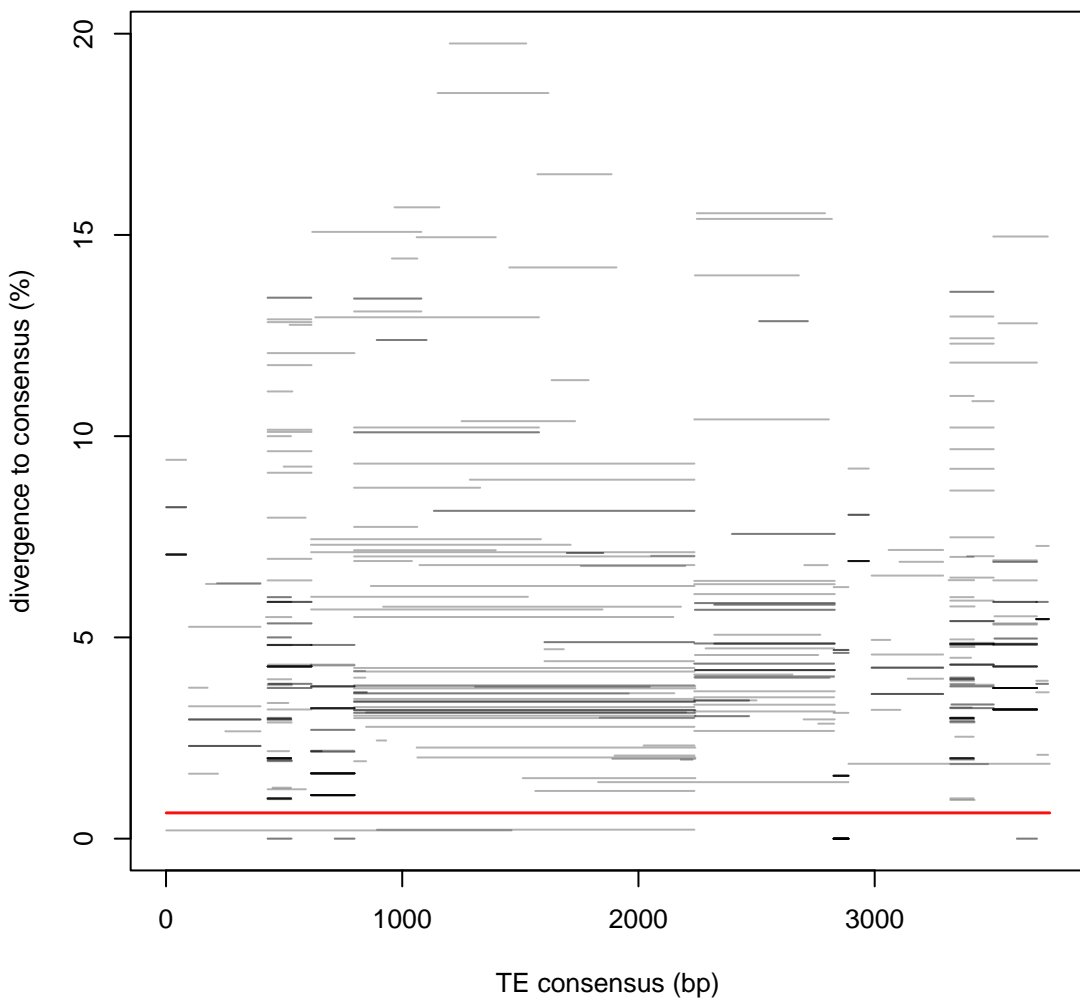
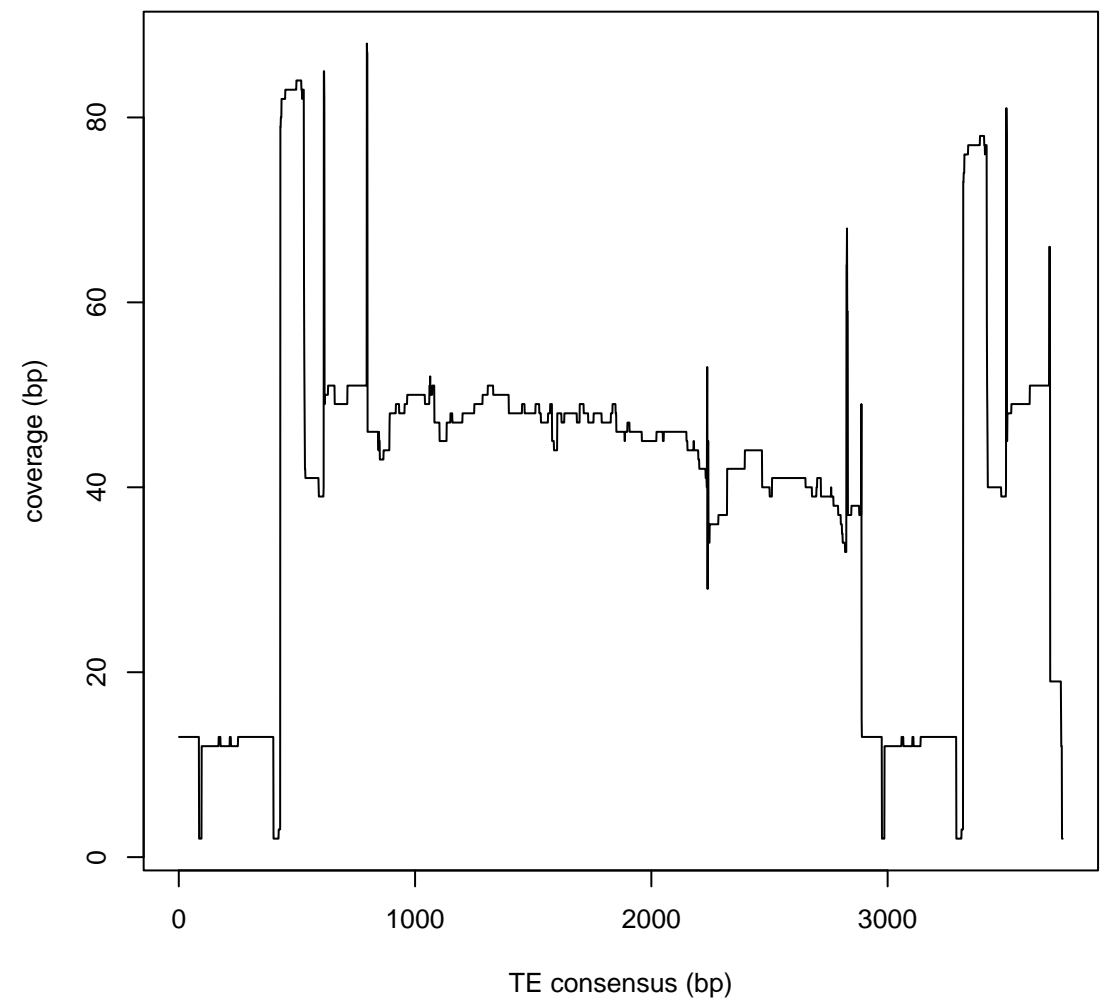


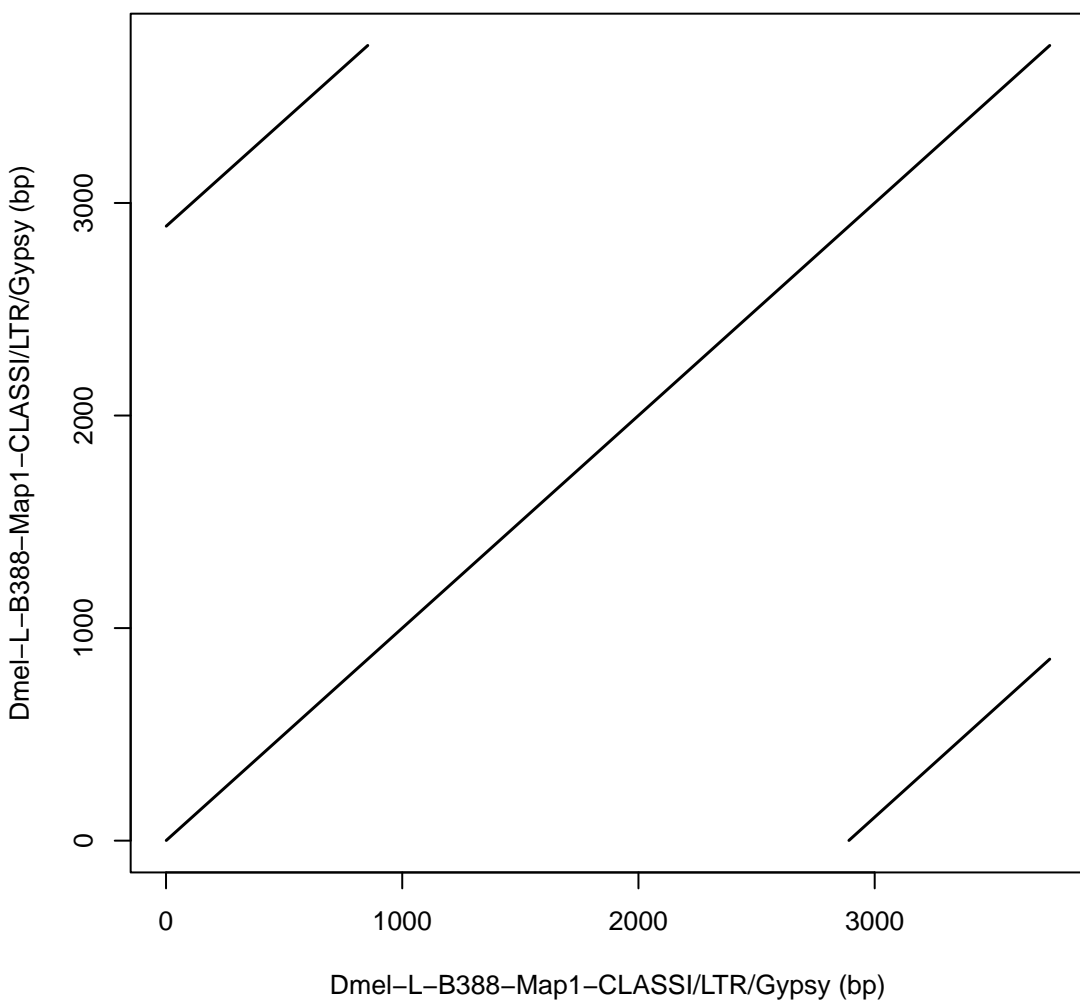
TE: Dmel-L-B388-Map1-CLASSI/LTR/Gypsy
consensus size: 3741bp; fragments: 485; full length: 1 (>=3366.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

