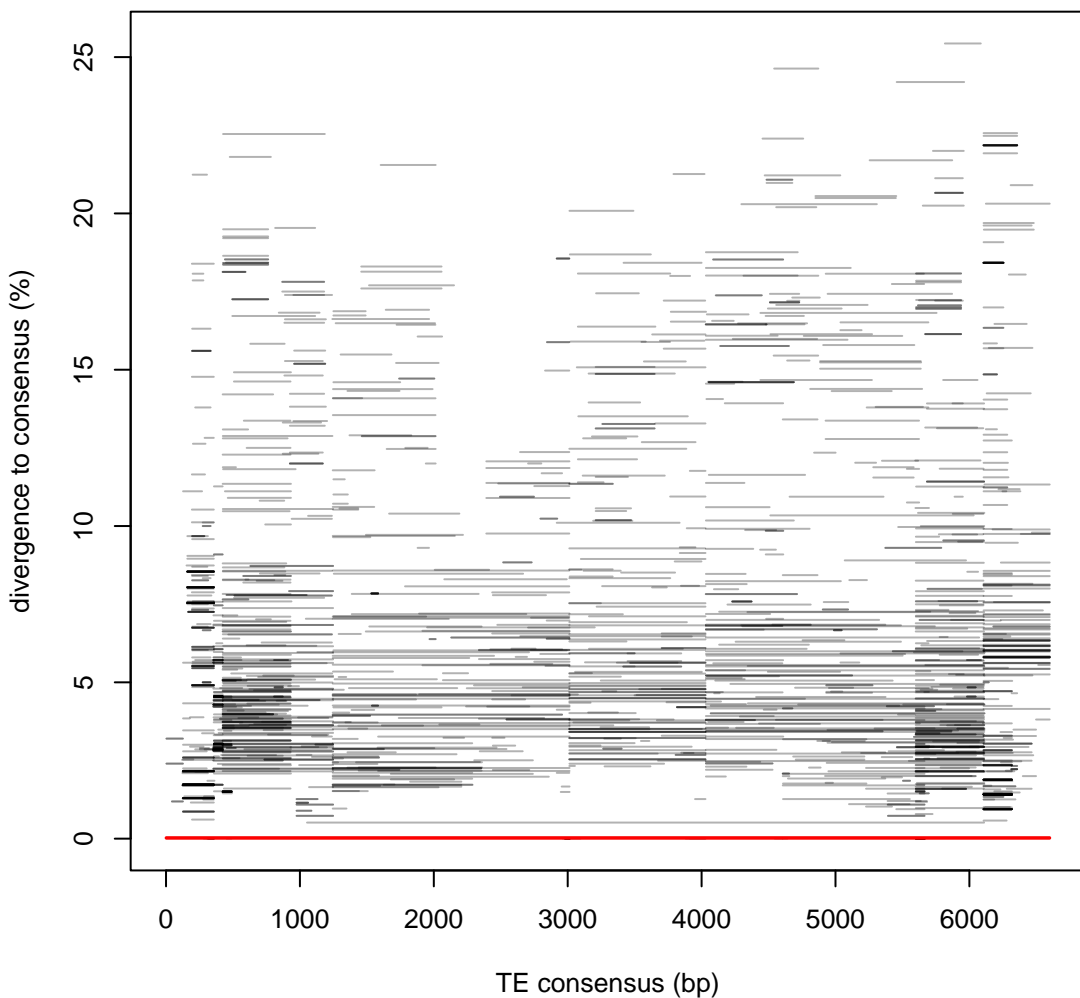
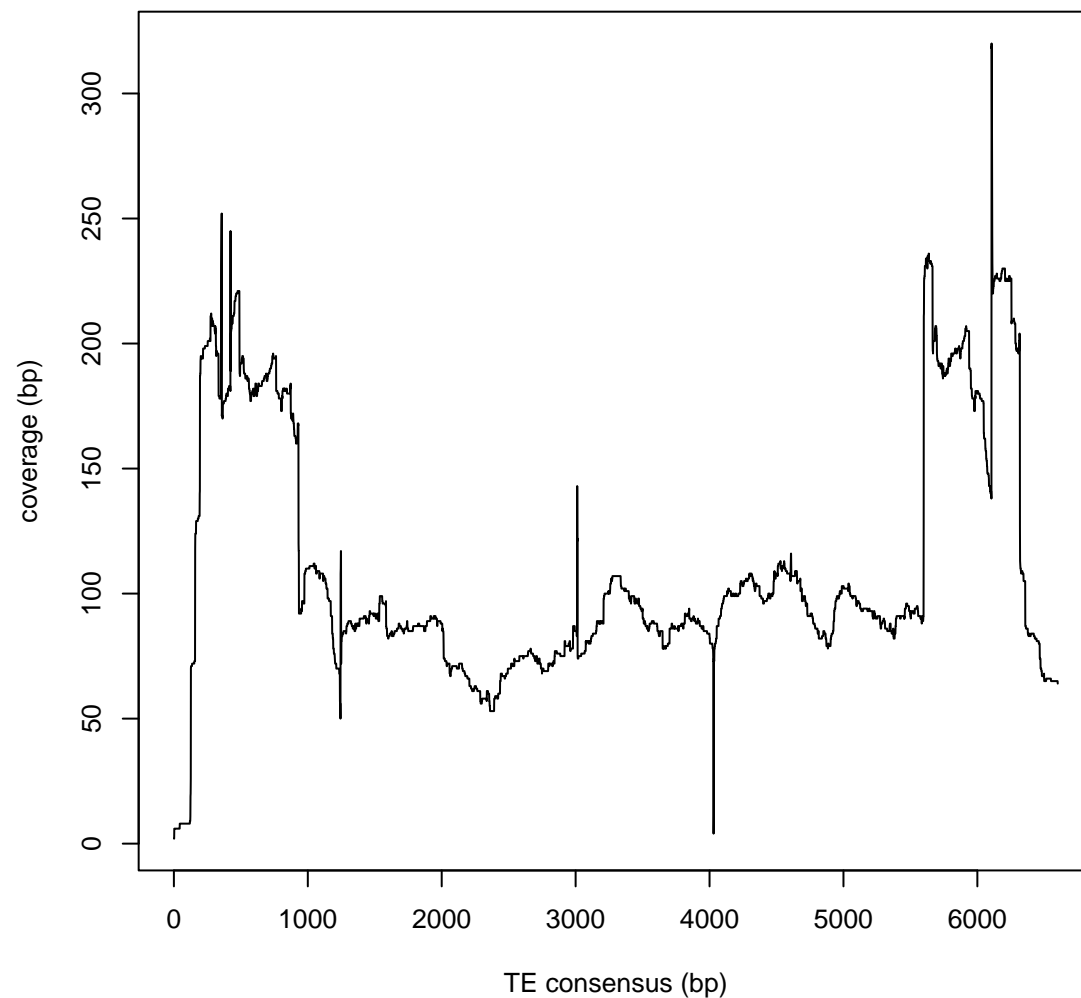


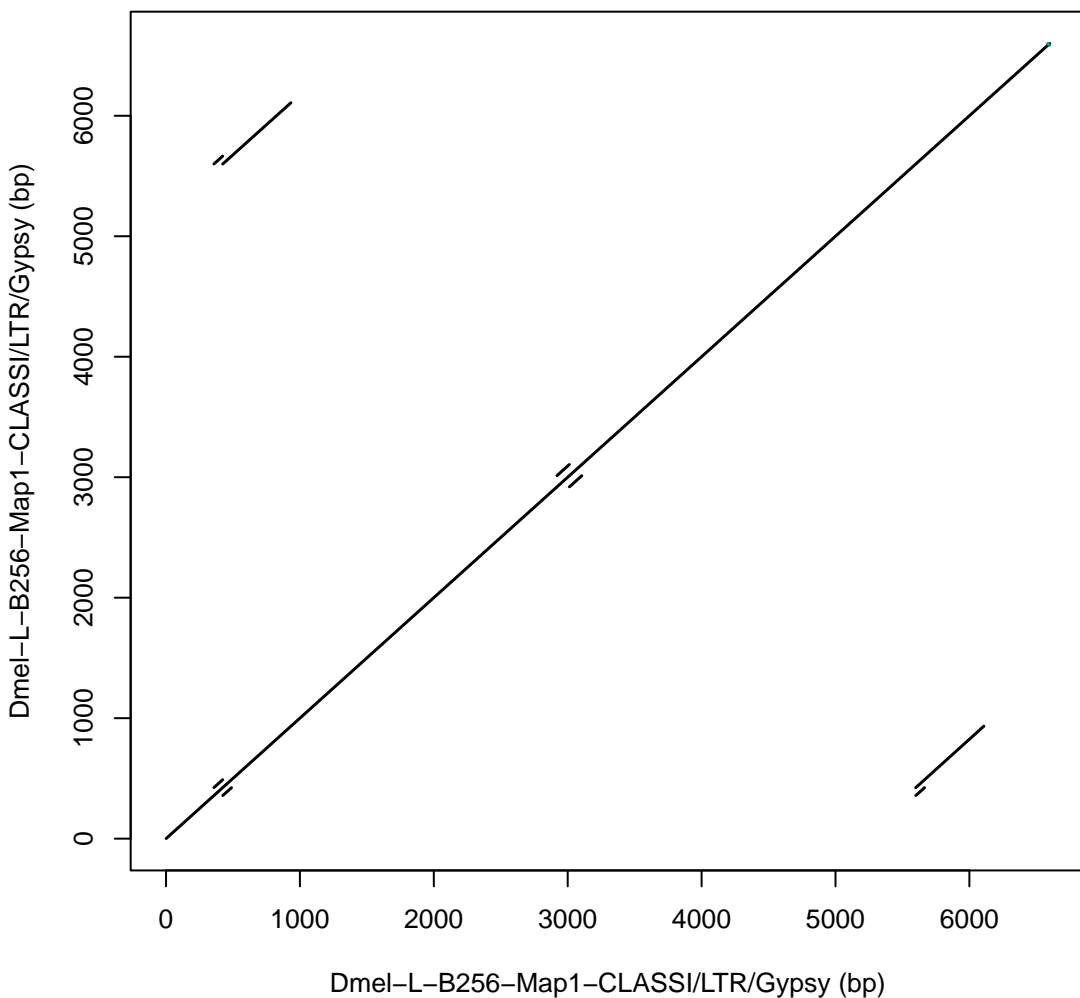
TE: Dmel-L-B256-Map1-CLASSI/LTR/Gypsy  
consensus size: 6600bp; fragments: 1842; full length: 2 (>=5940bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

