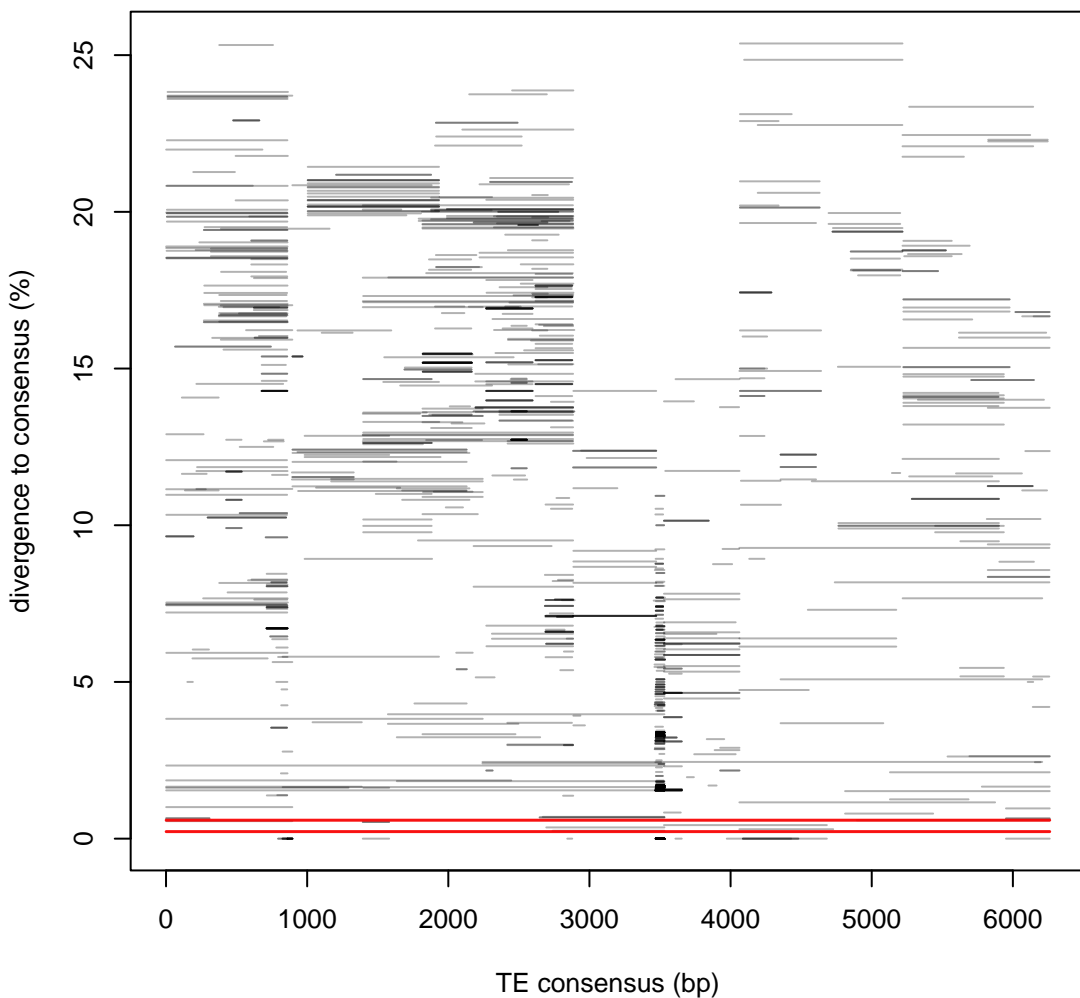
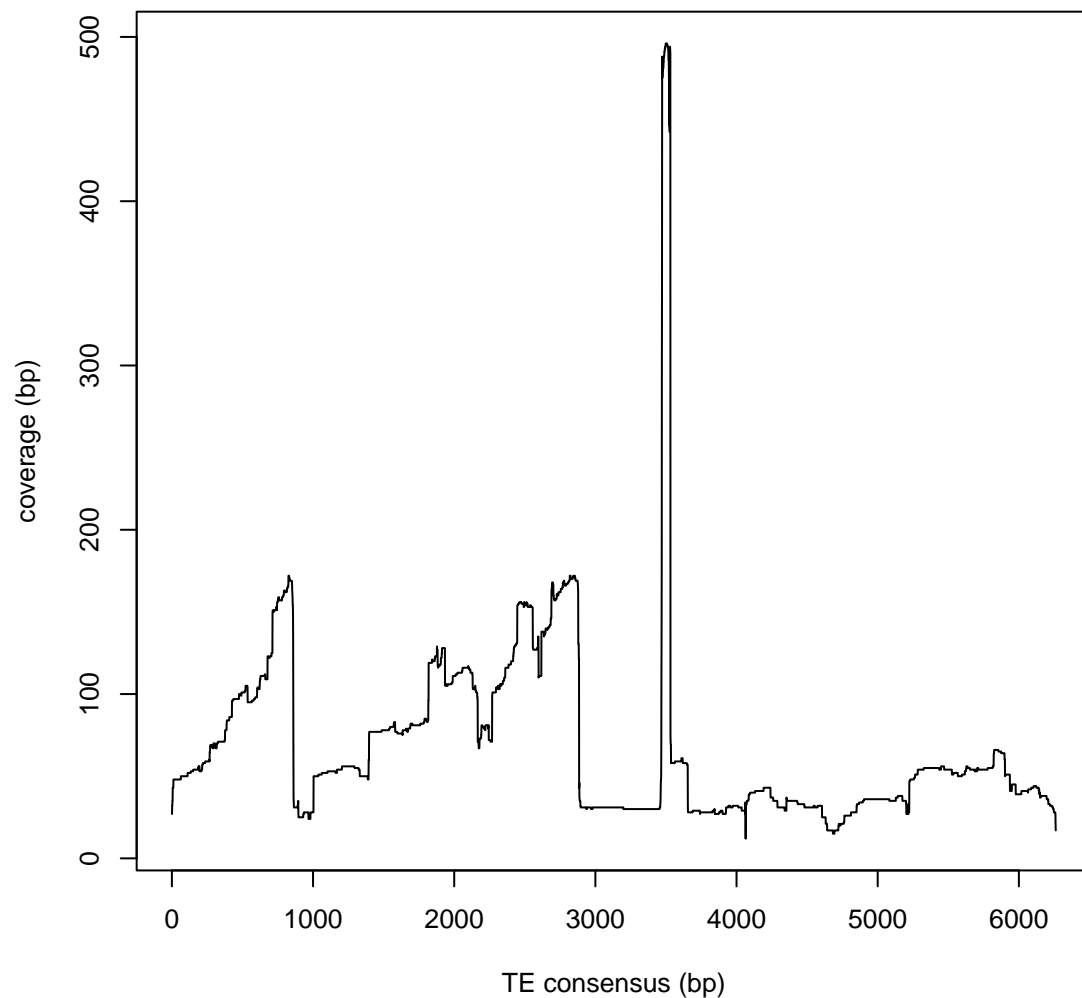


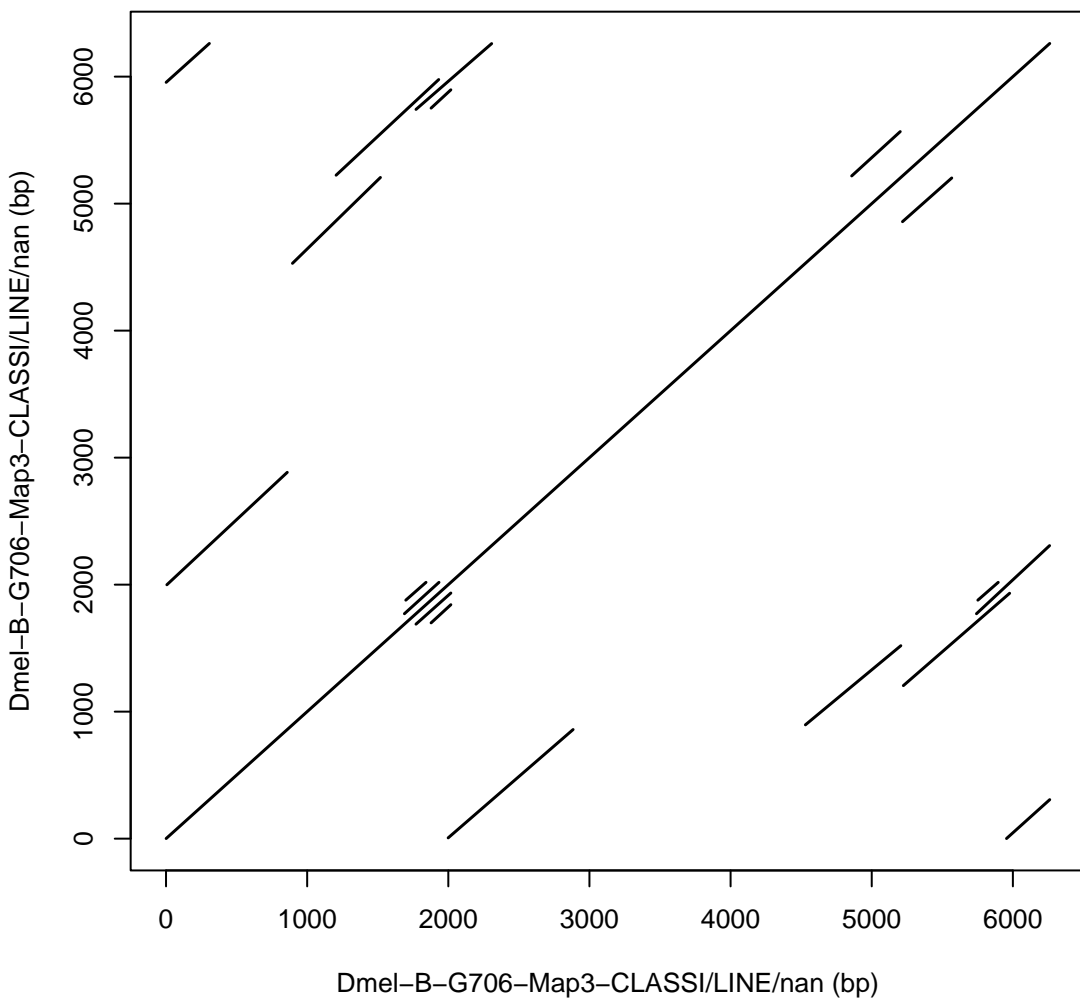
TE: Dmel-B-G706-Map3-CLASSI/LINE/nan  
consensus size: 6261bp; fragments: 1320; full length: 2 ( $\geq 5634.9$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

