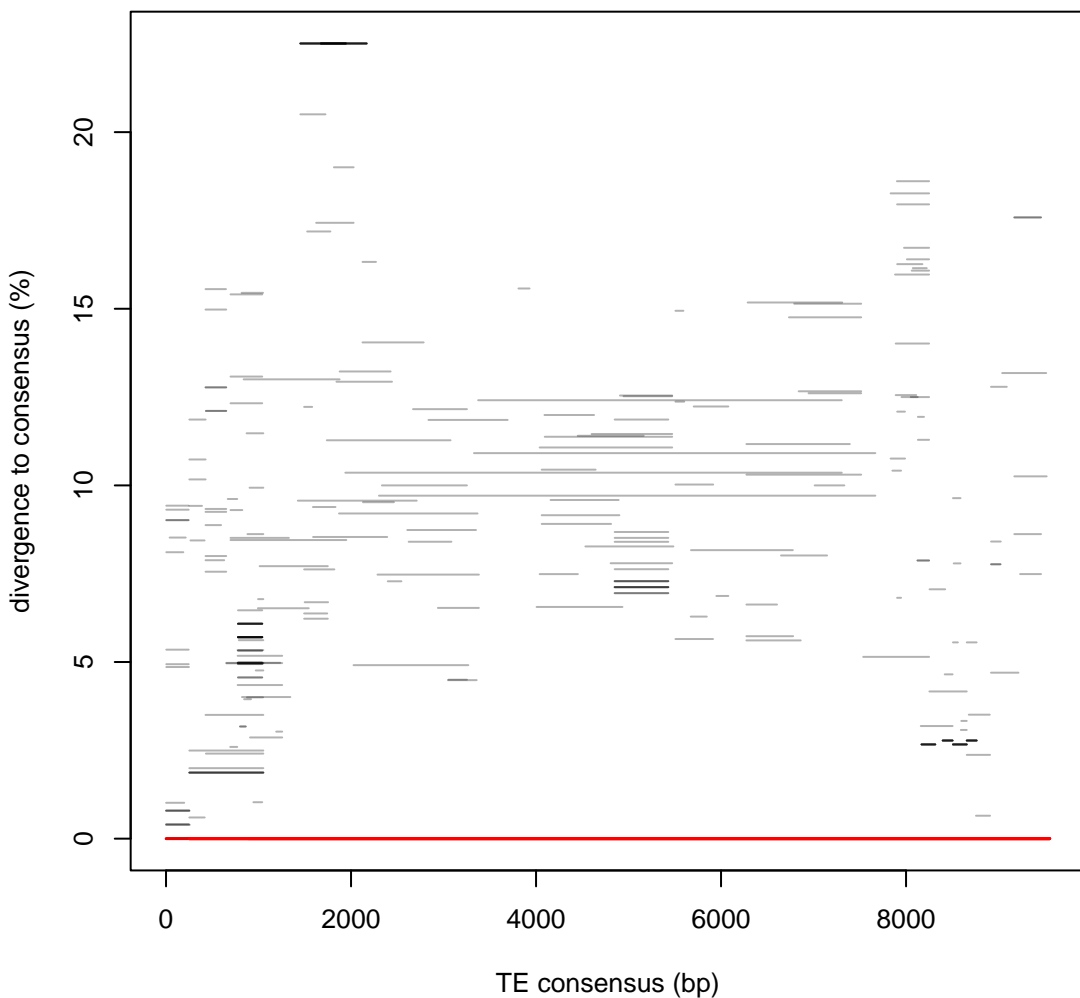
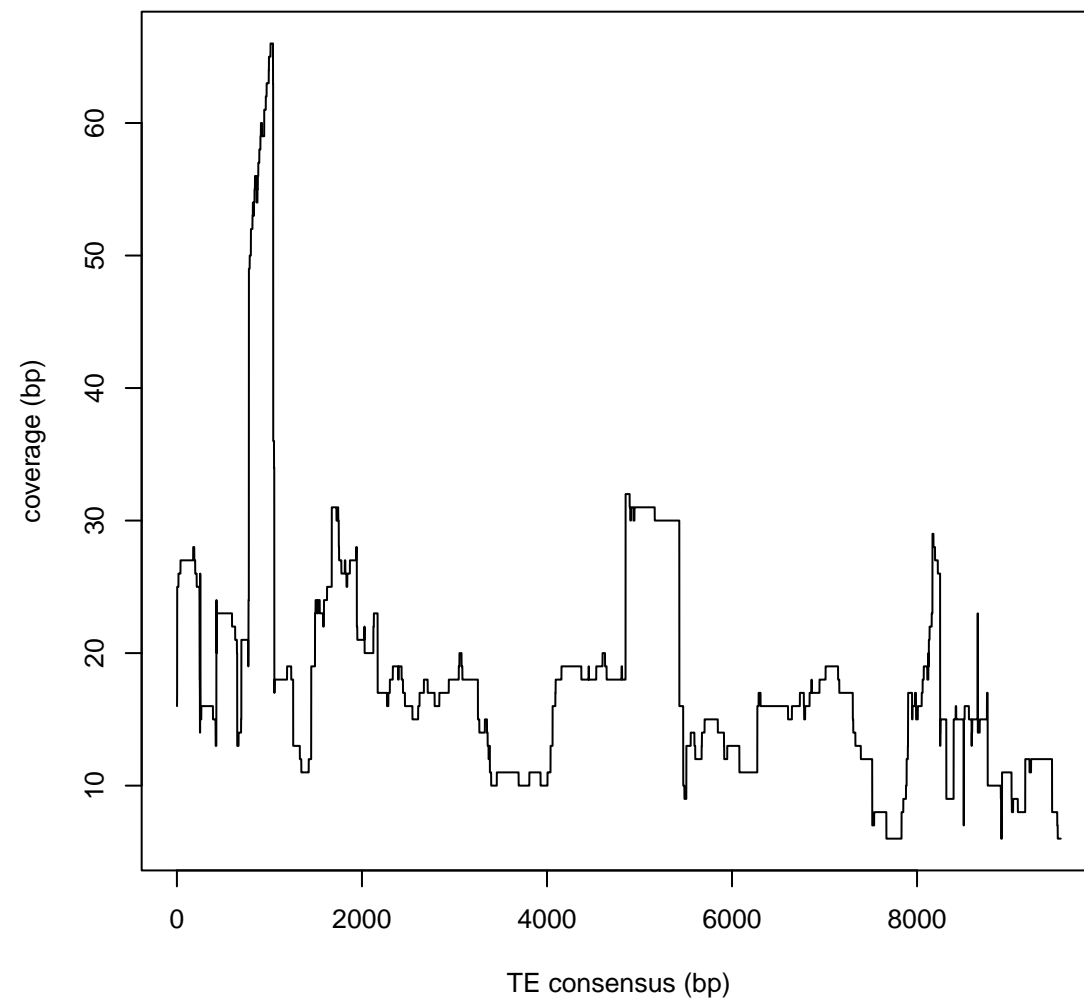


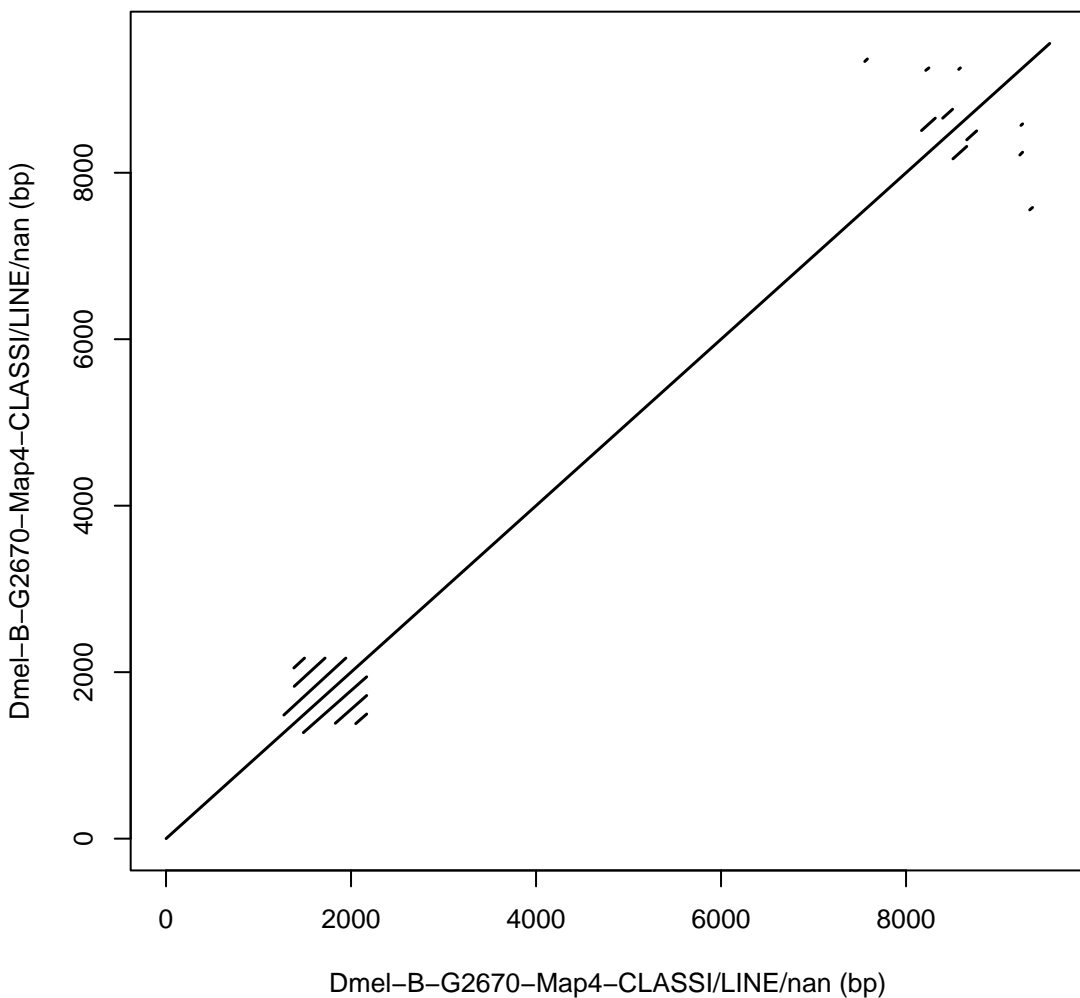
TE: Dmel-B-G2670-Map4-CLASSI/LINE/nan  
consensus size: 9553bp; fragments: 291; full length: 4 (>=8597.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

