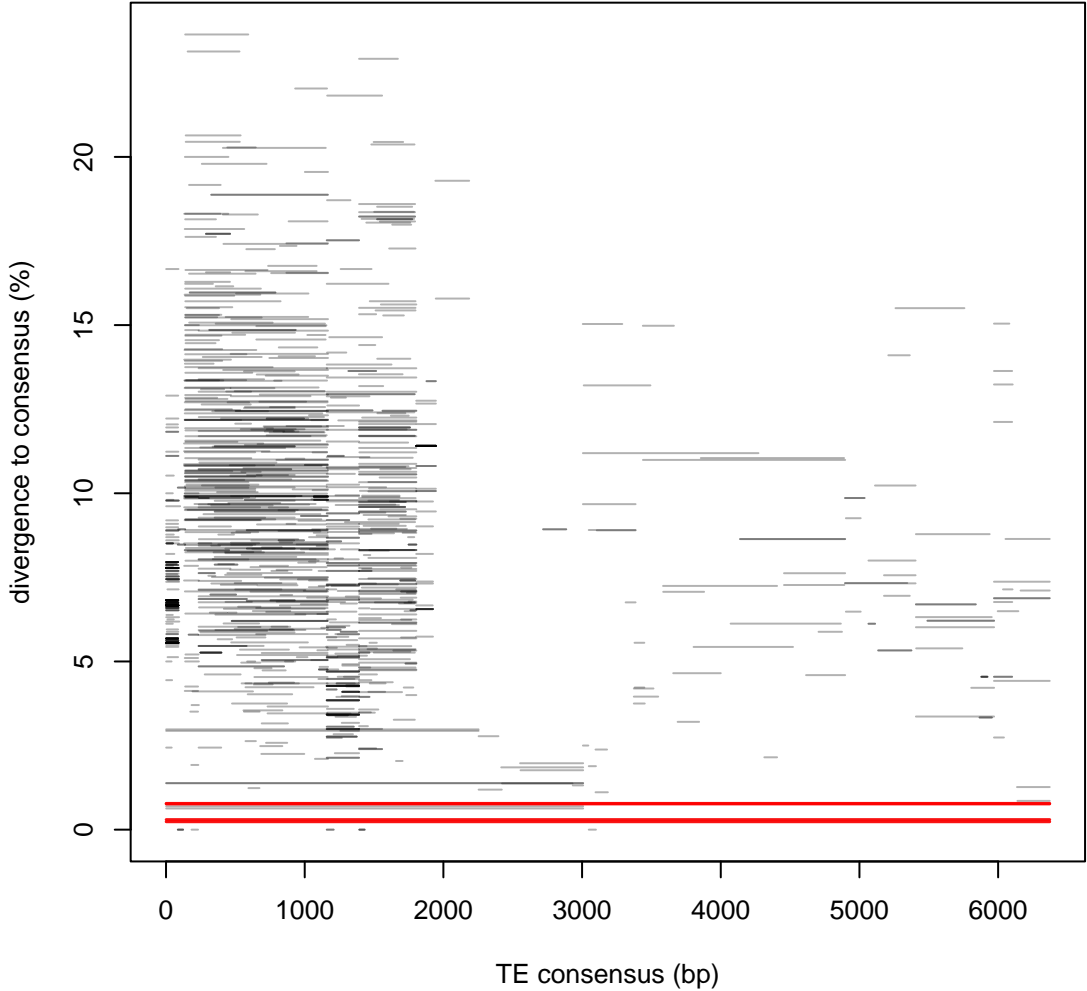
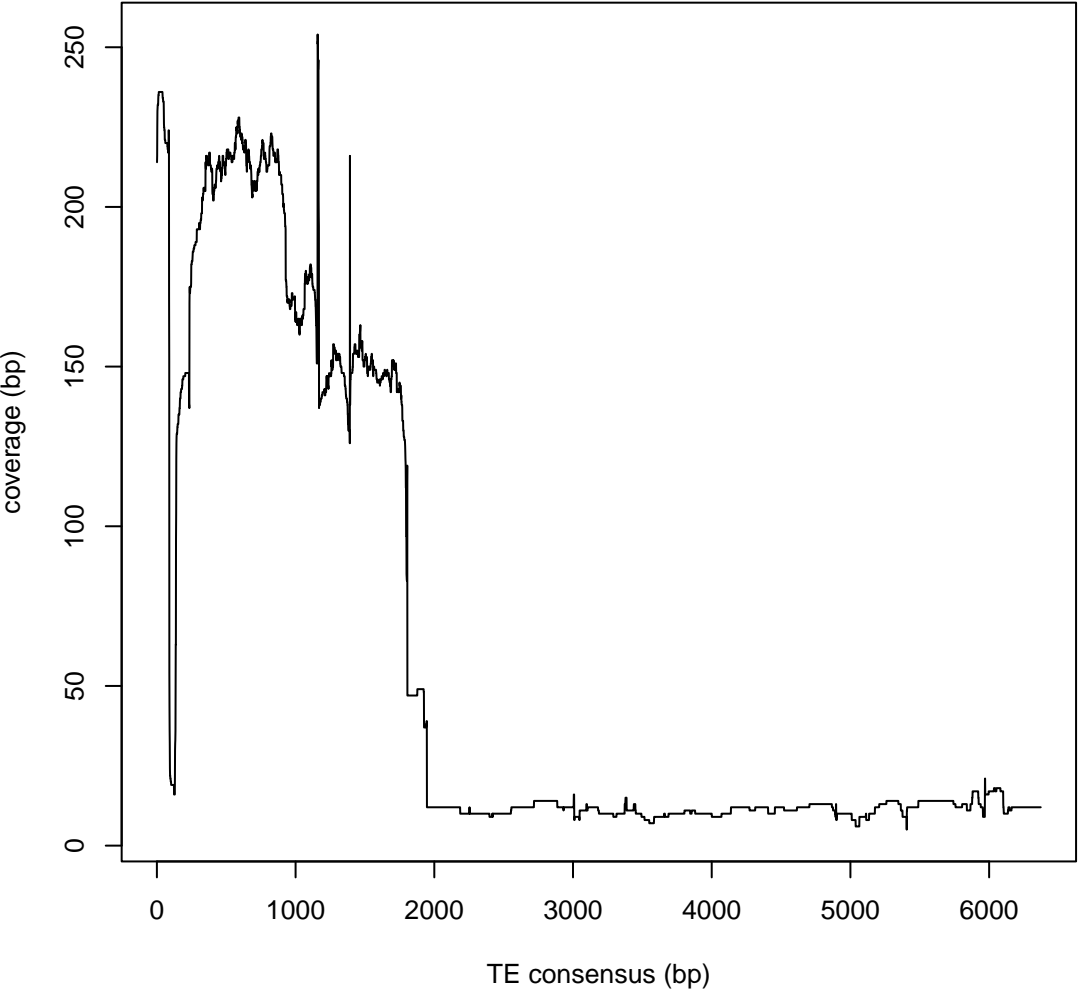


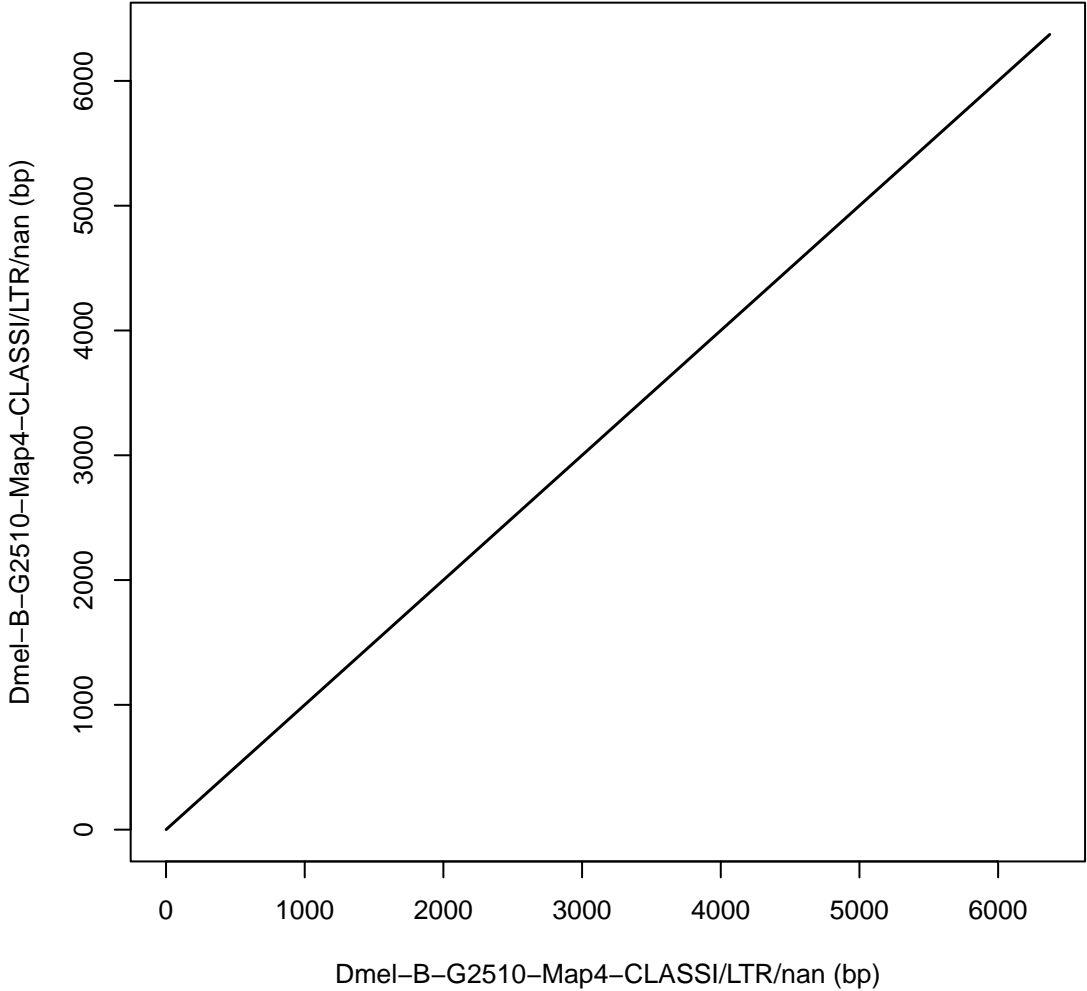
TE: Dmel-B-G2510-Map4-CLASSI/LTR/nan  
consensus size: 6372bp; fragments: 1194; full length: 4 (>=5734.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

