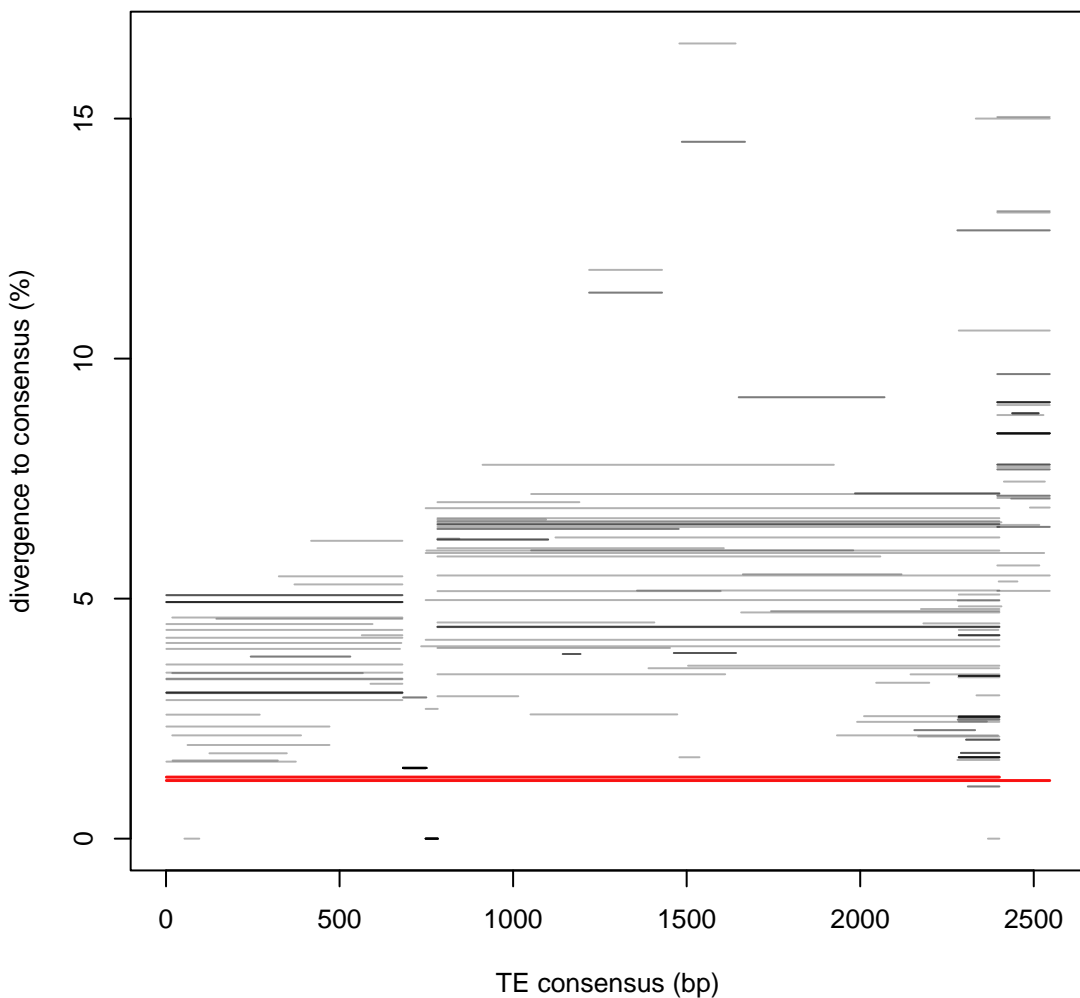
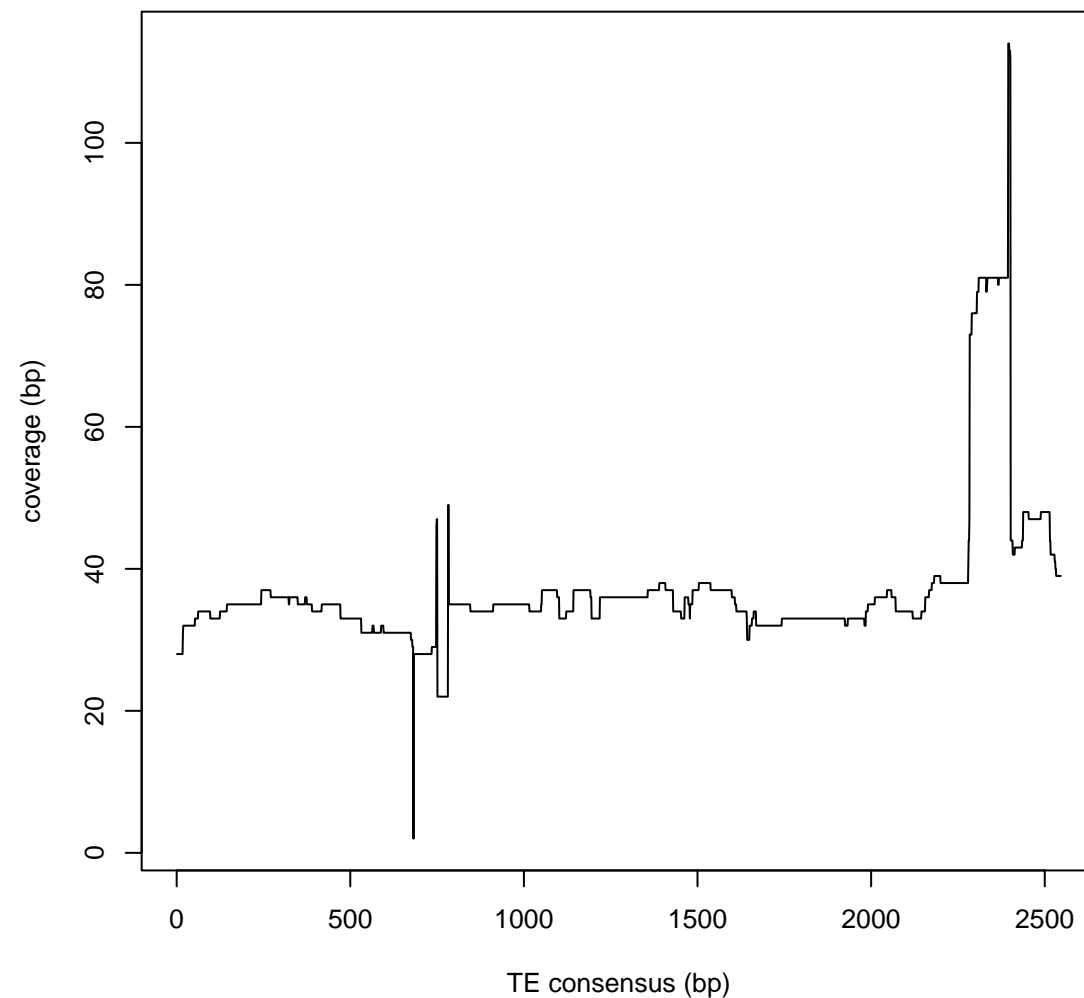


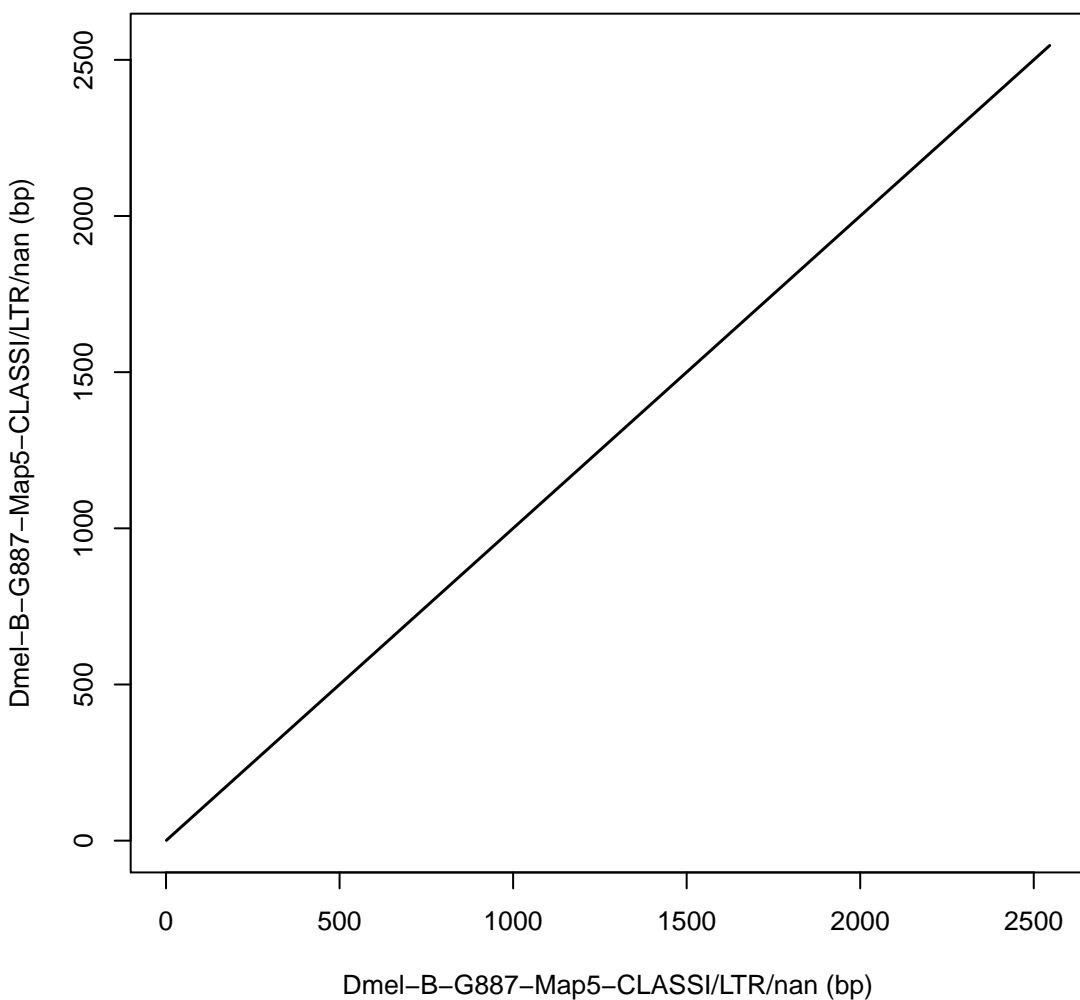
TE: Dmel-B-G887-Map5-CLASSI/LTR/nan  
consensus size: 2546bp; fragments: 241; full length: 2 ( $\geq 2291.4$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

