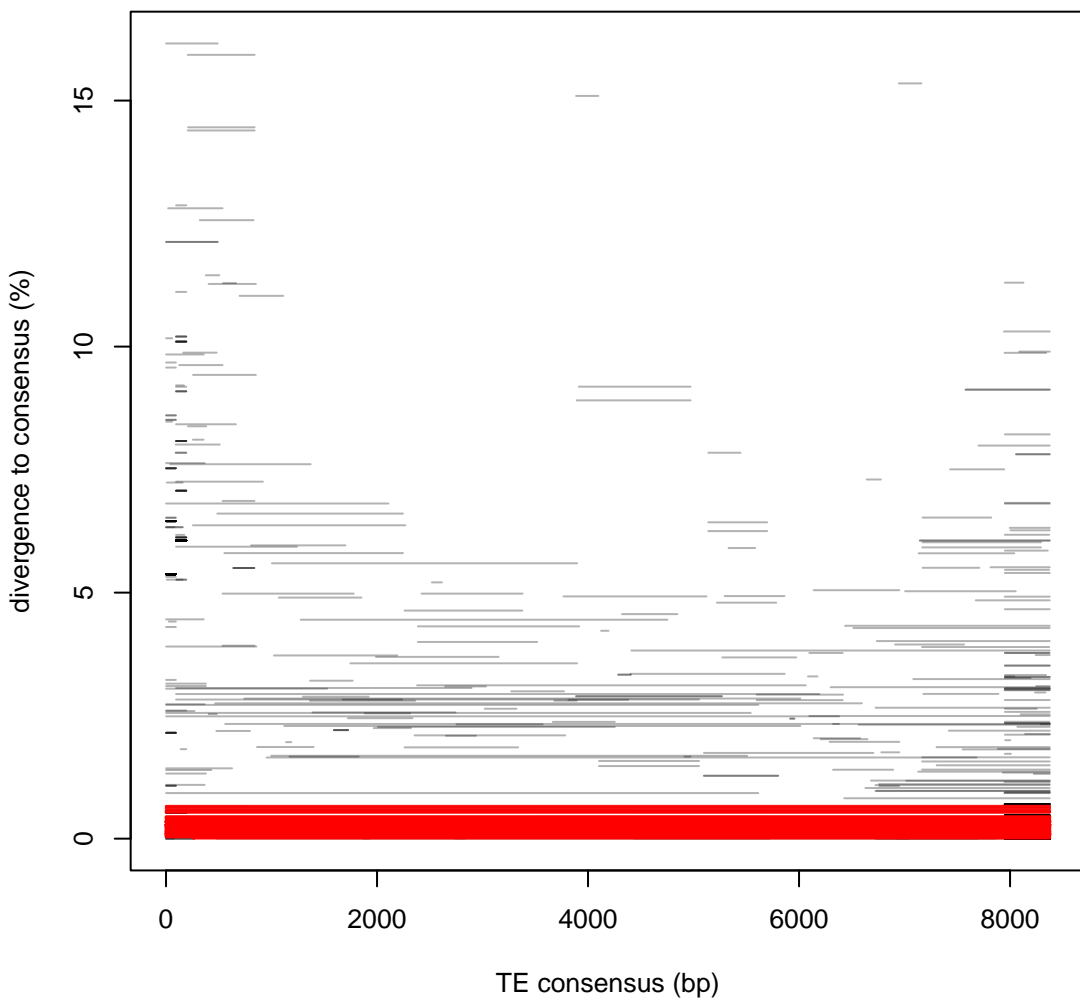
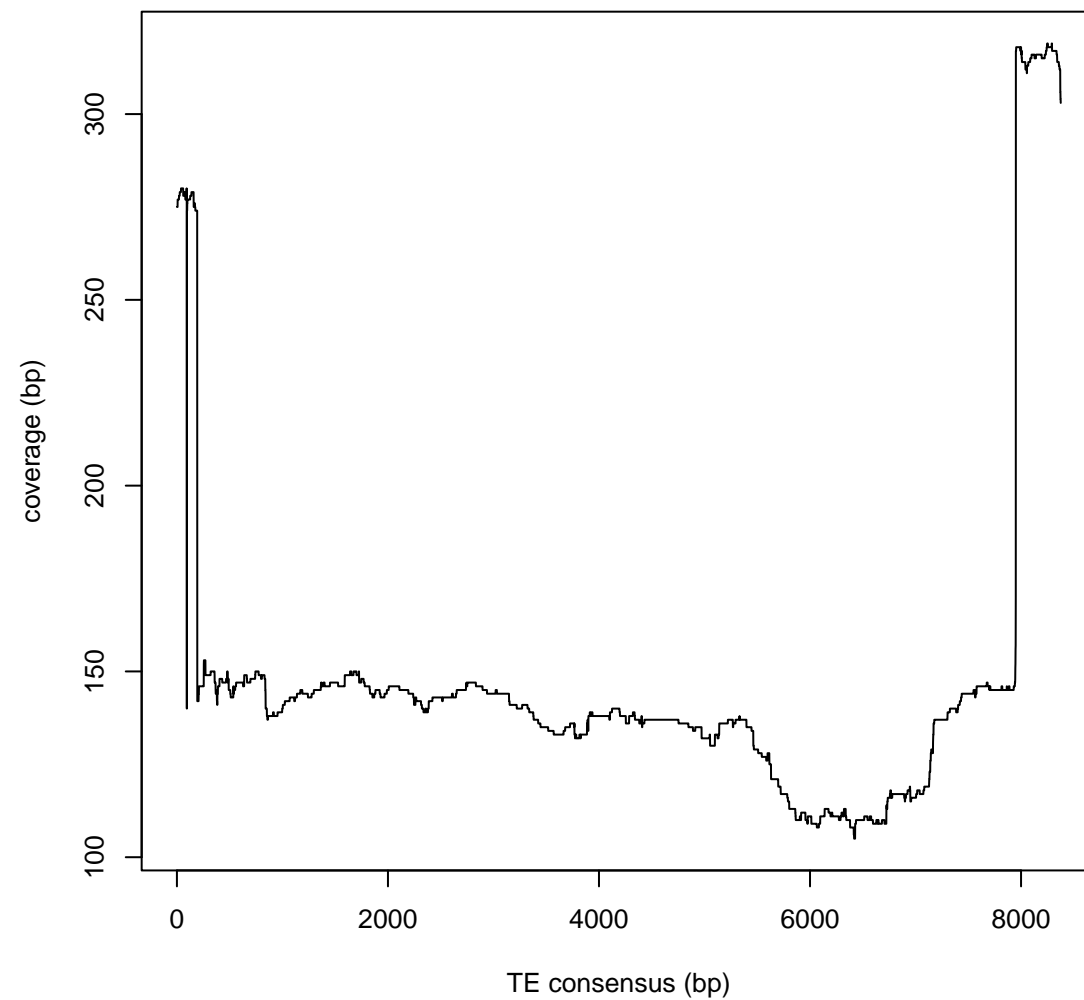


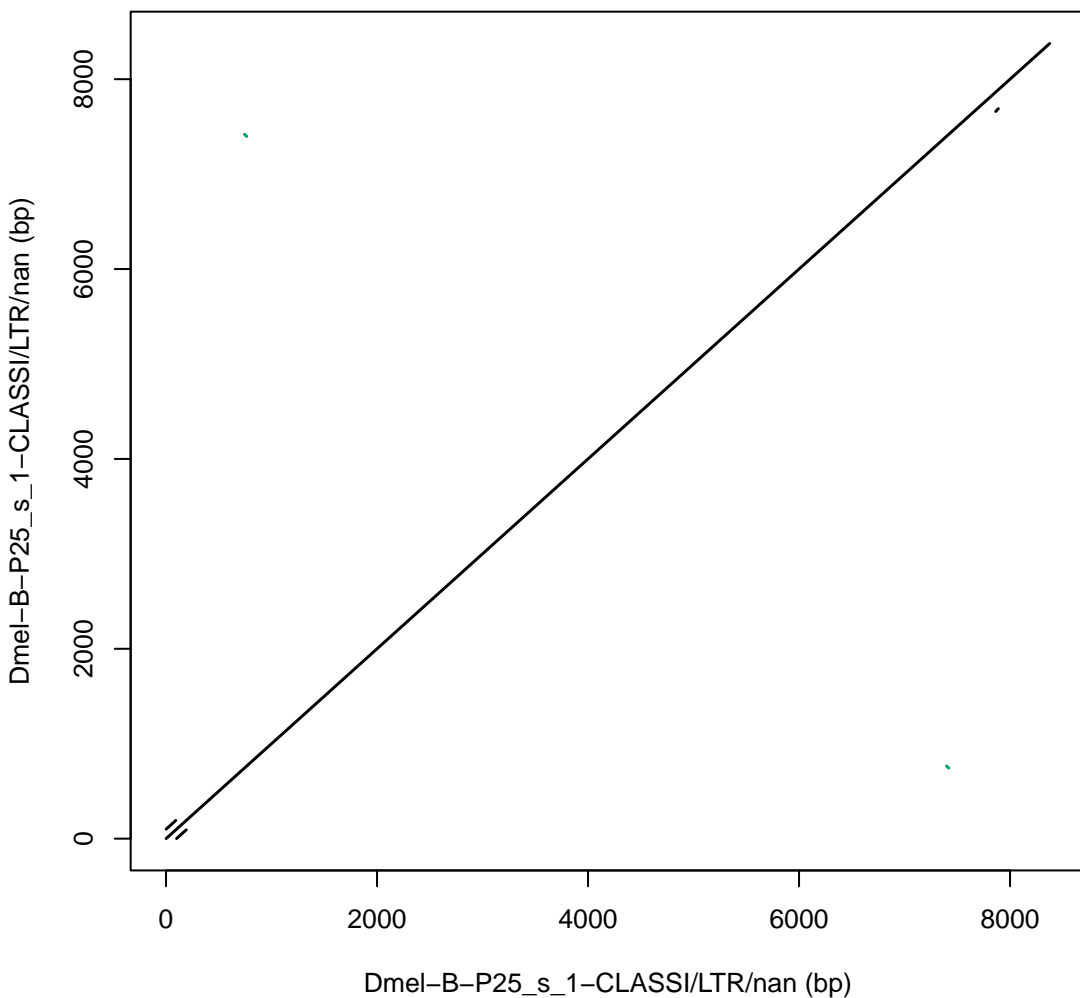
TE: Dmel-B-P25\_s\_1-CLASSI/LTR/nan  
consensus size: 8376bp; fragments: 816; full length: 72 ( $\geq 7538.4$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

