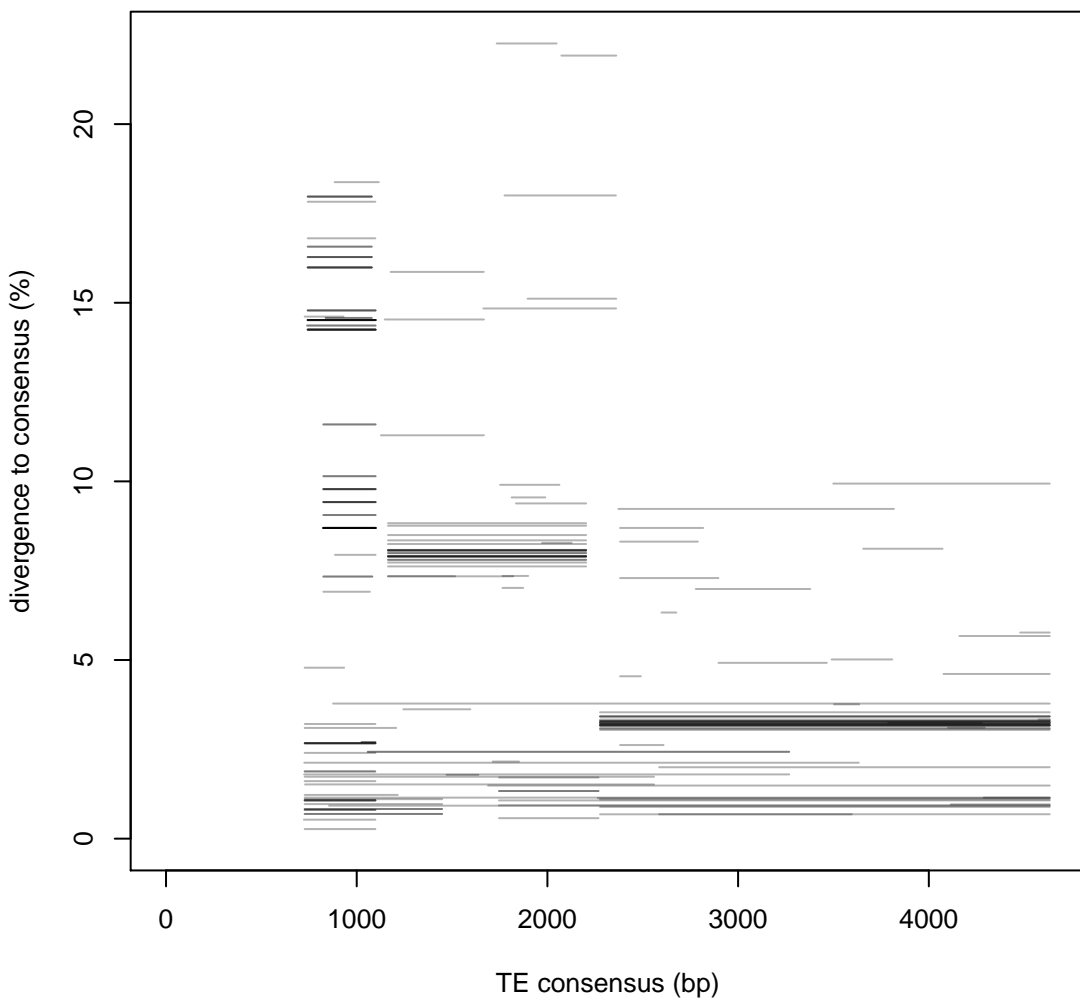
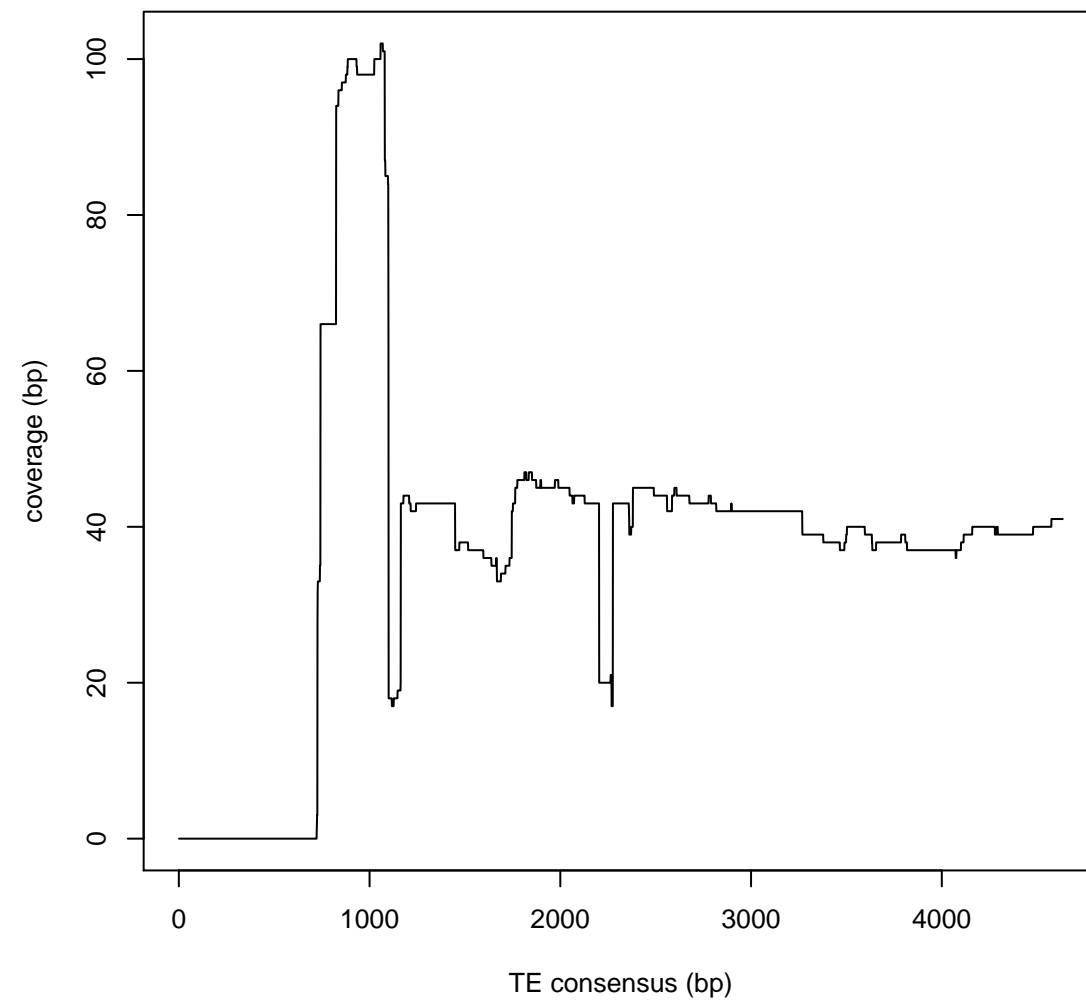


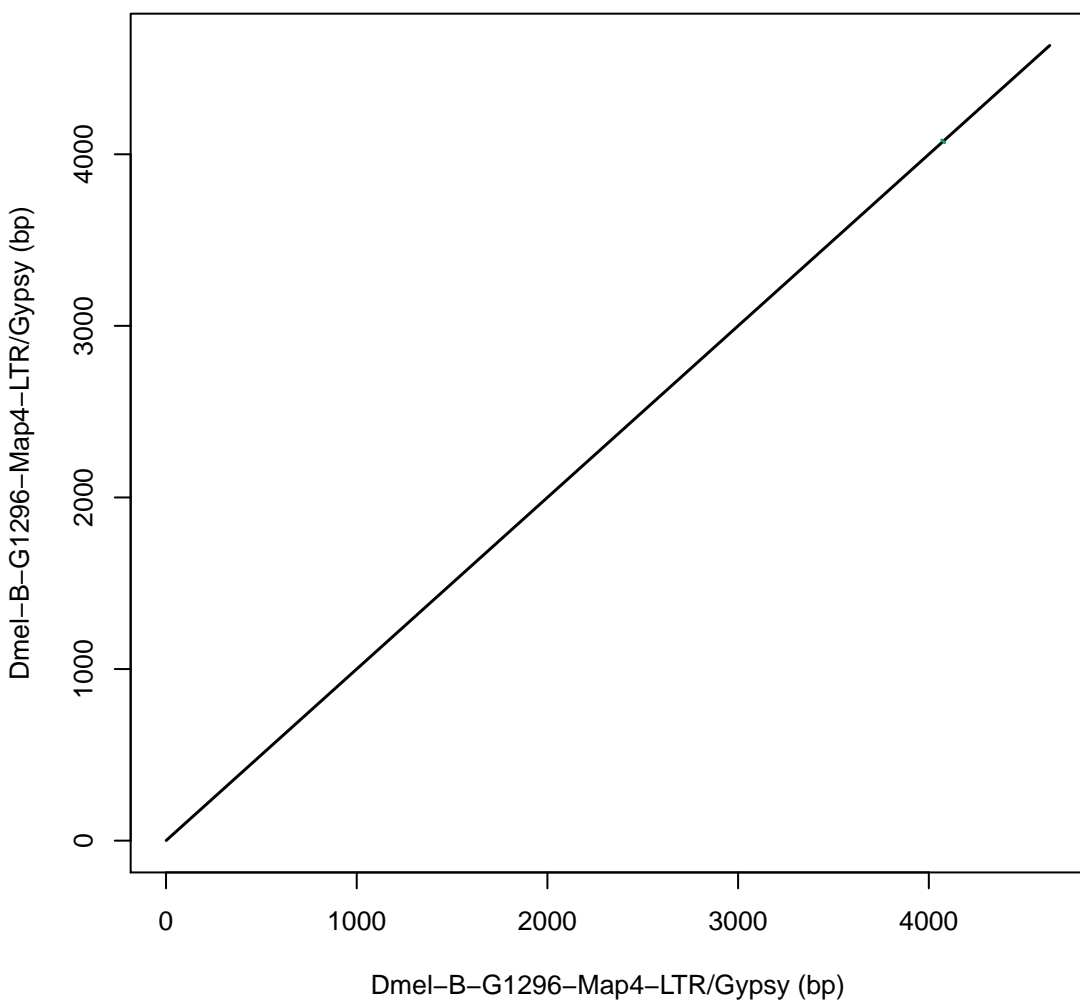
TE: Dmel-B-G1296-Map4-LTR/Gypsy  
consensus size: 4634bp; fragments: 202; full length: 0 ( $\geq 4170.6$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

