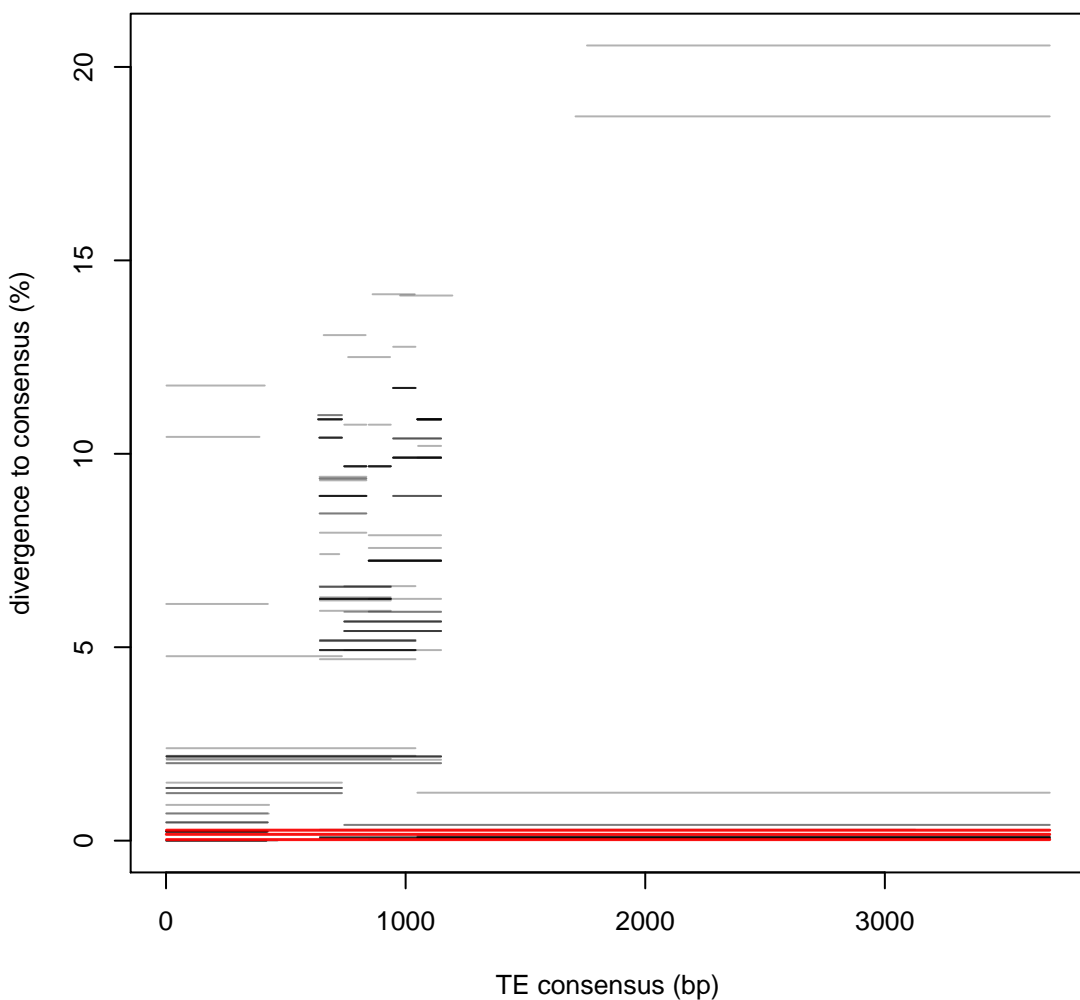
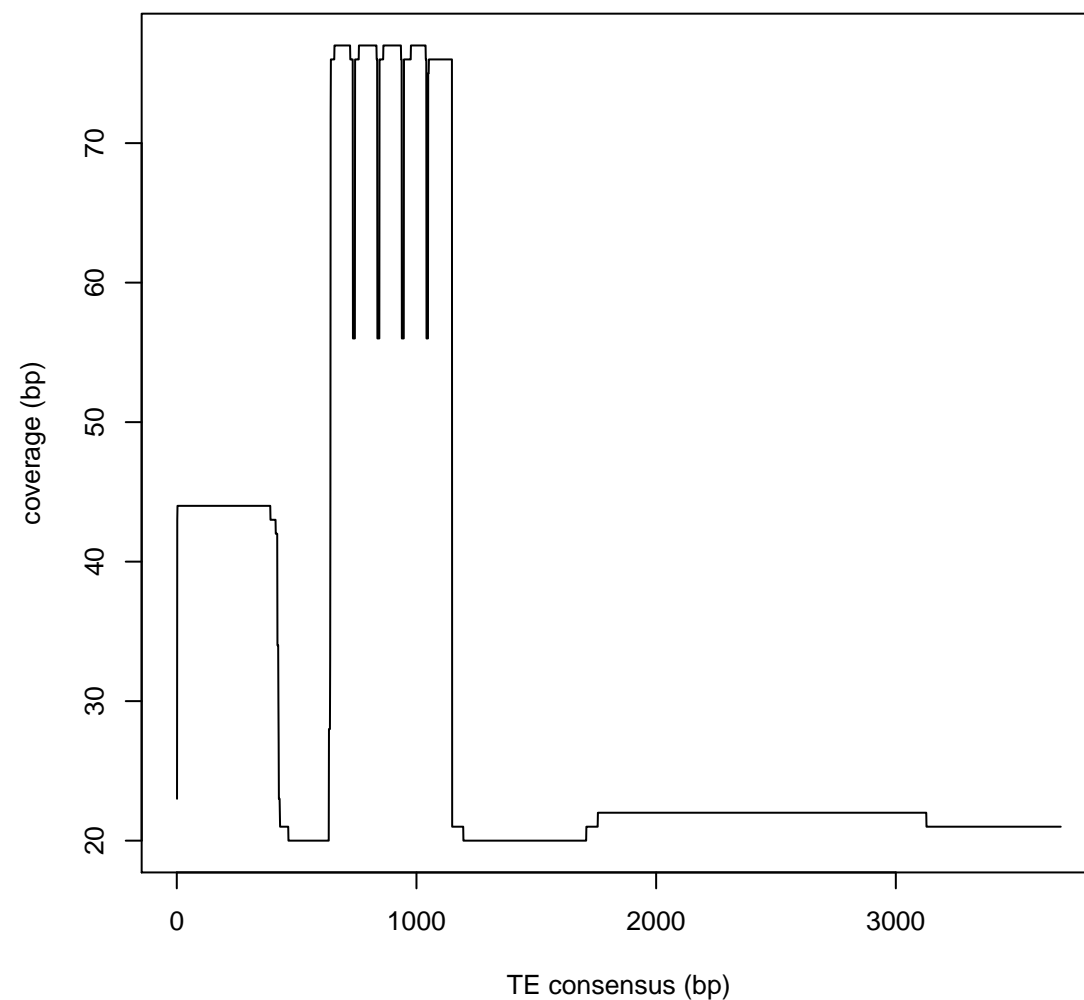


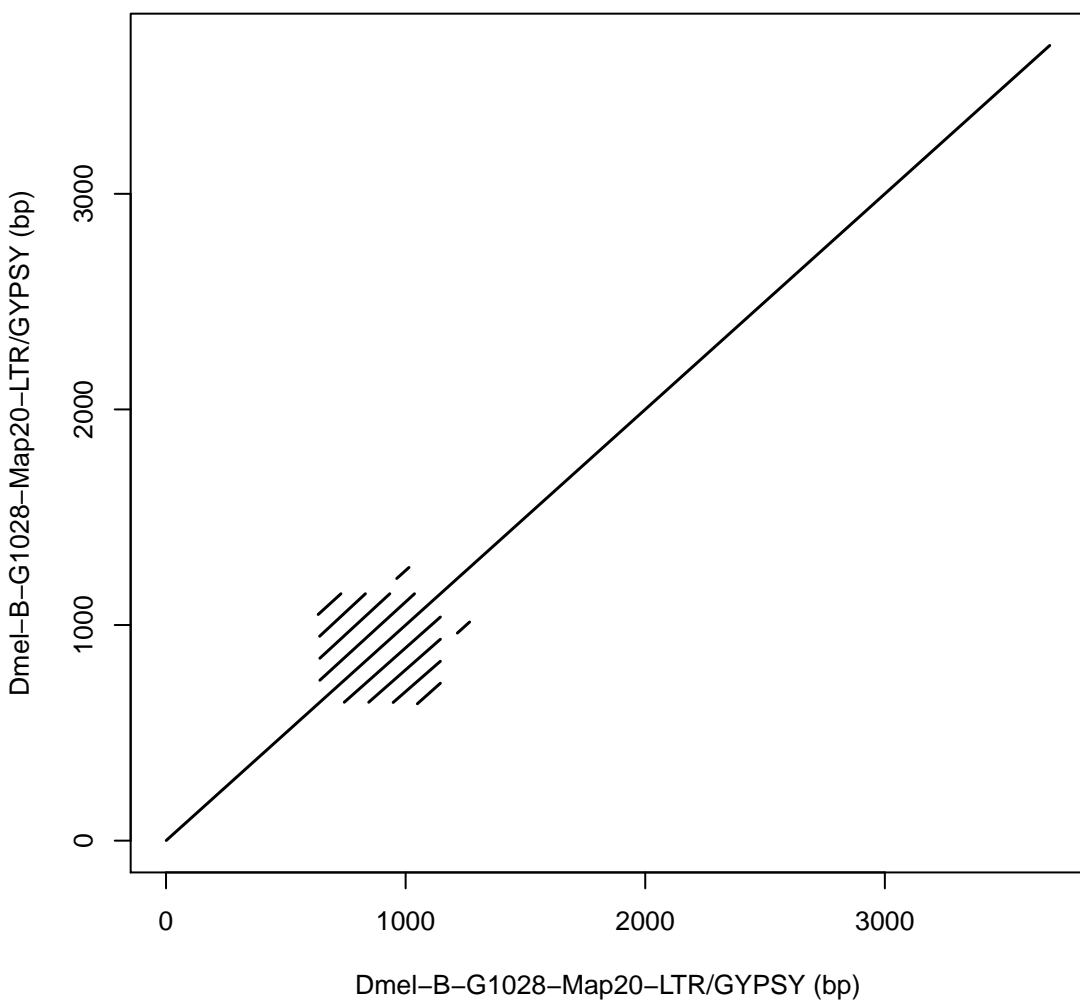
TE: Dmel-B-G1028-Map20-LTR/GYPSY  
consensus size: 3688bp; fragments: 186; full length: 3 (>=3319.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

