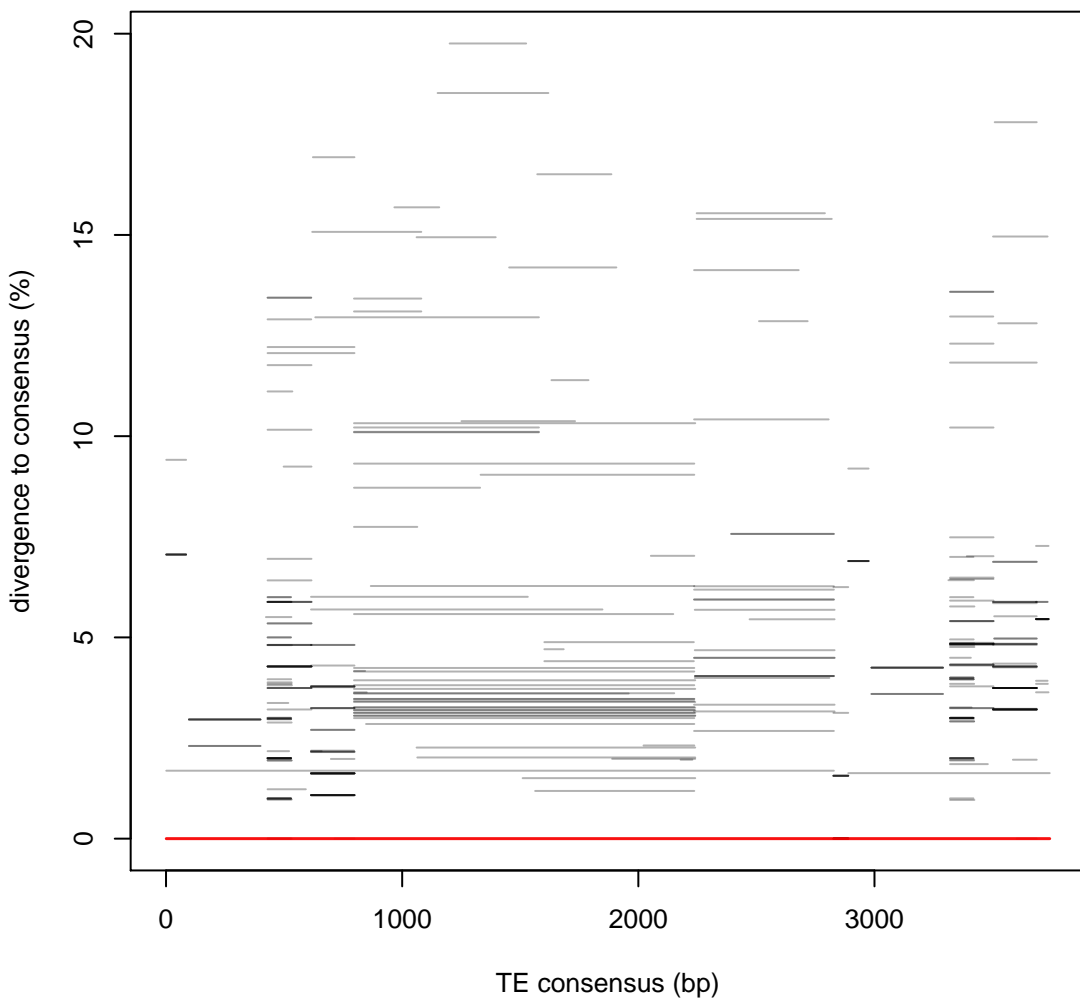
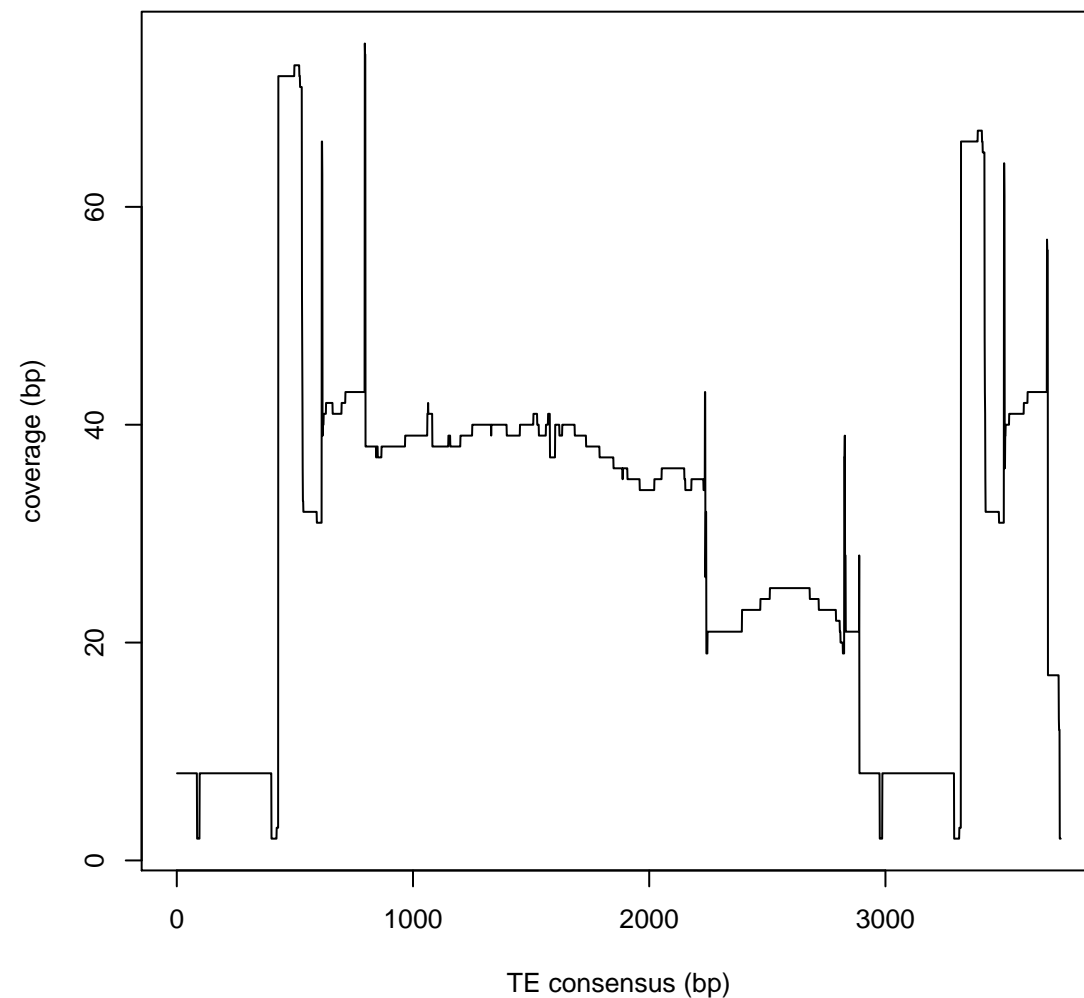


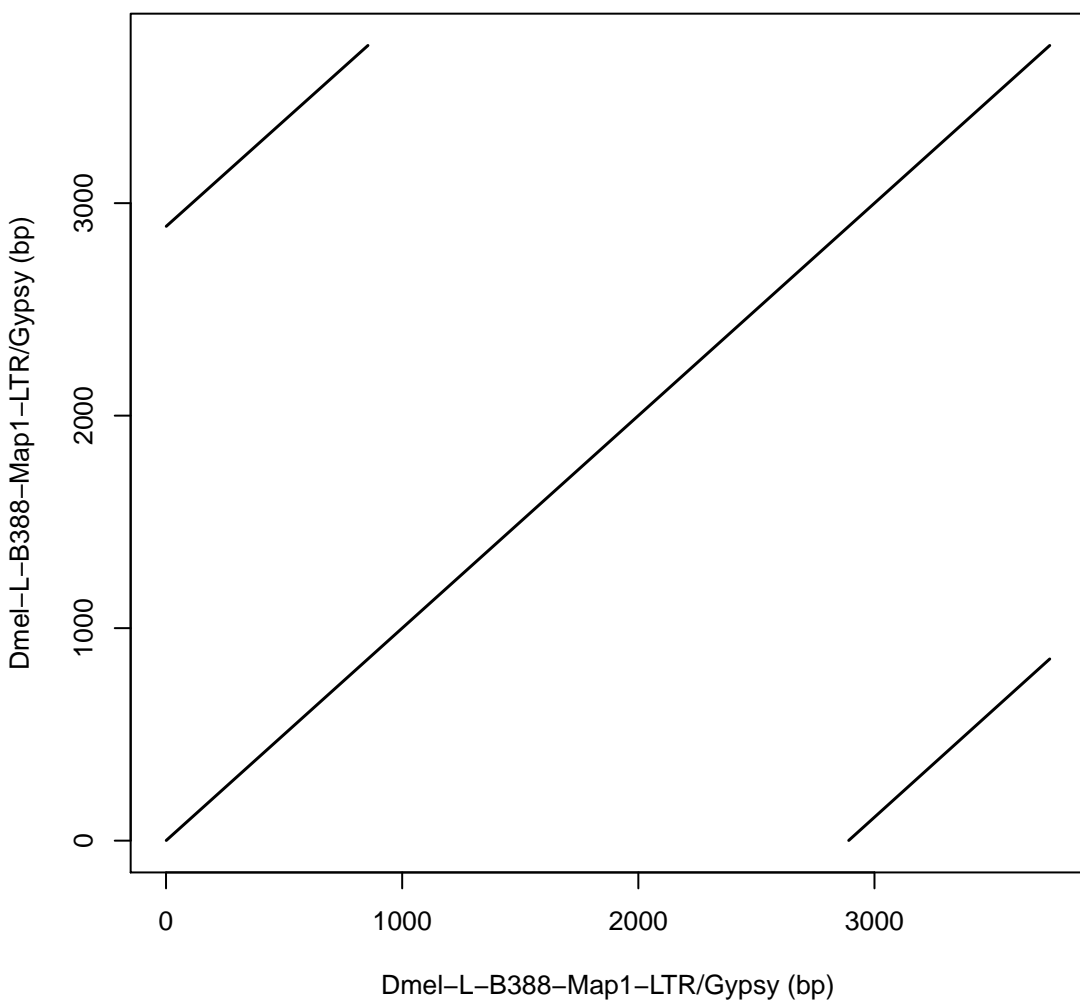
TE: Dmel-L-B388-Map1-LTR/Gypsy  
consensus size: 3742bp; fragments: 354; full length: 1 (>=3367.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

