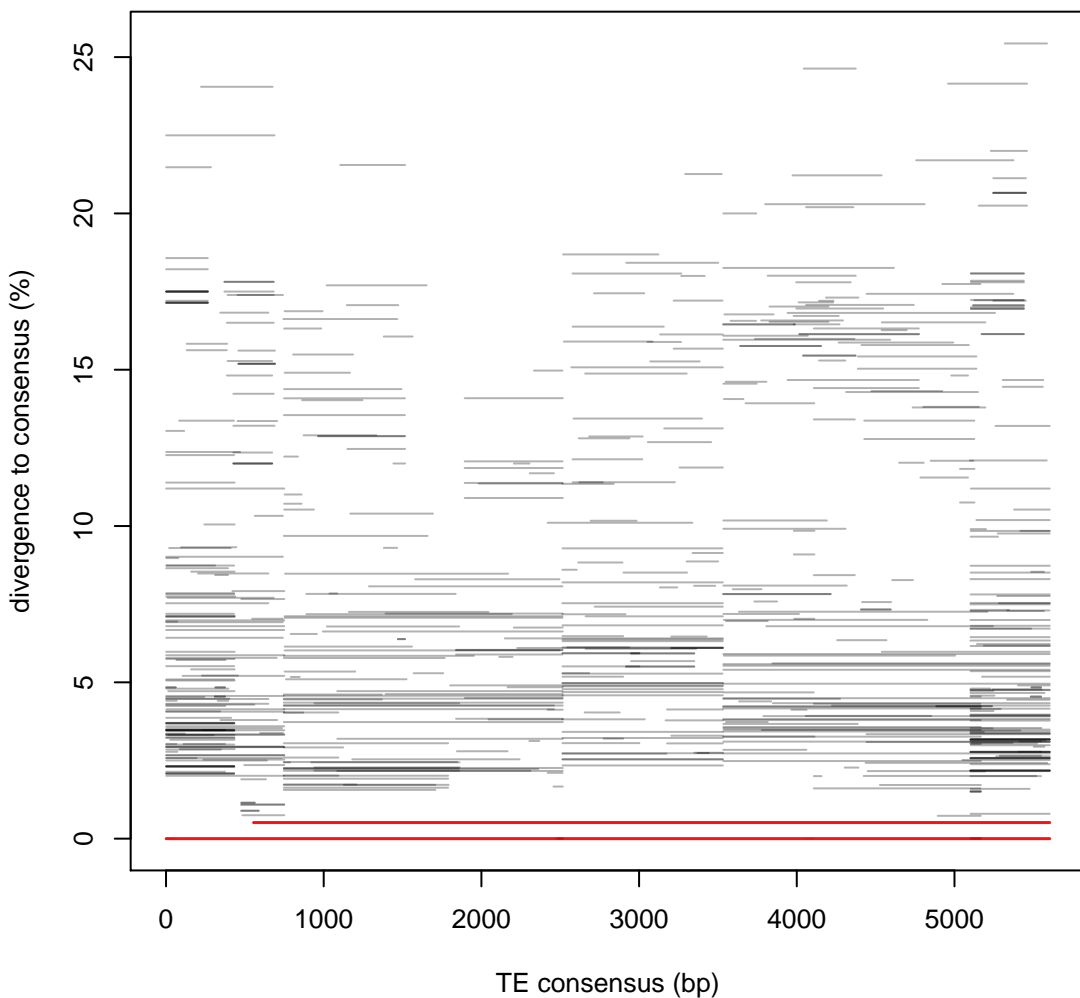
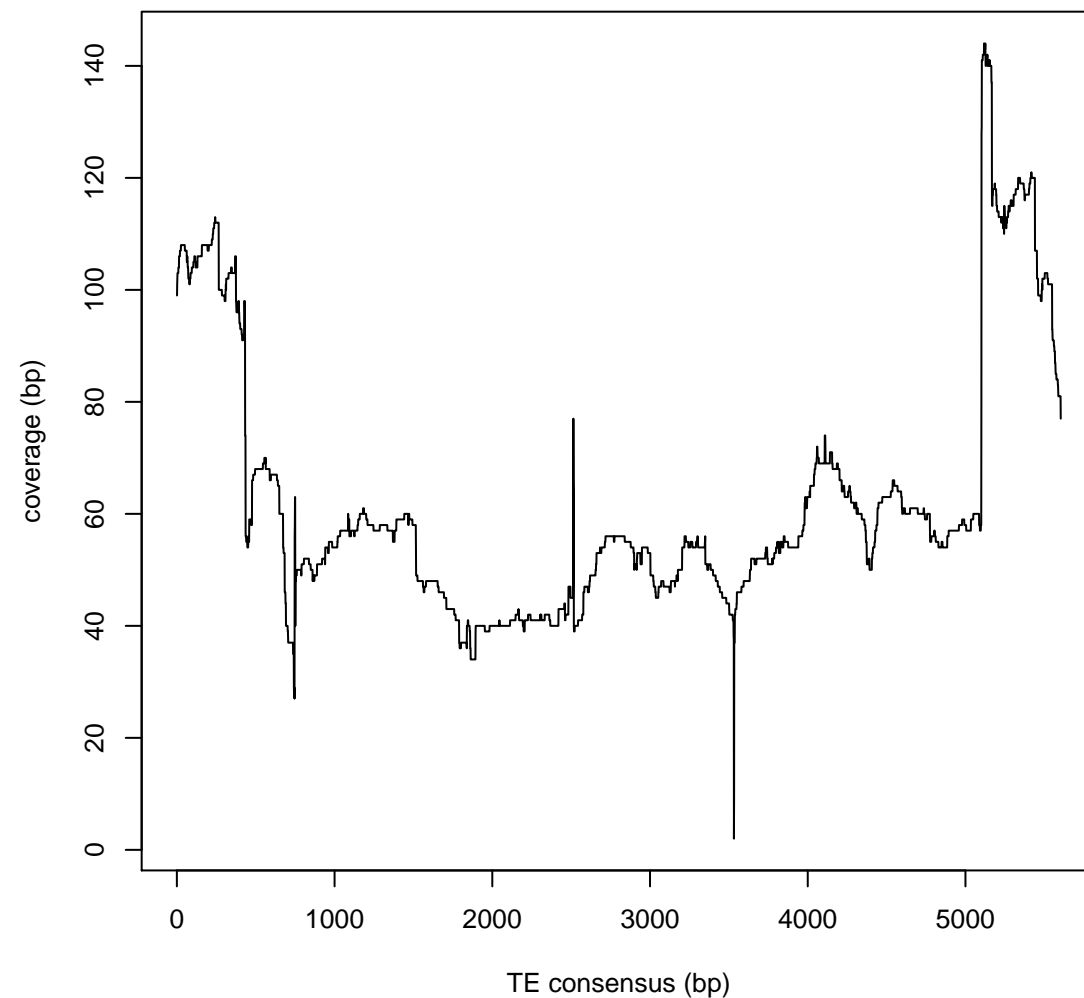


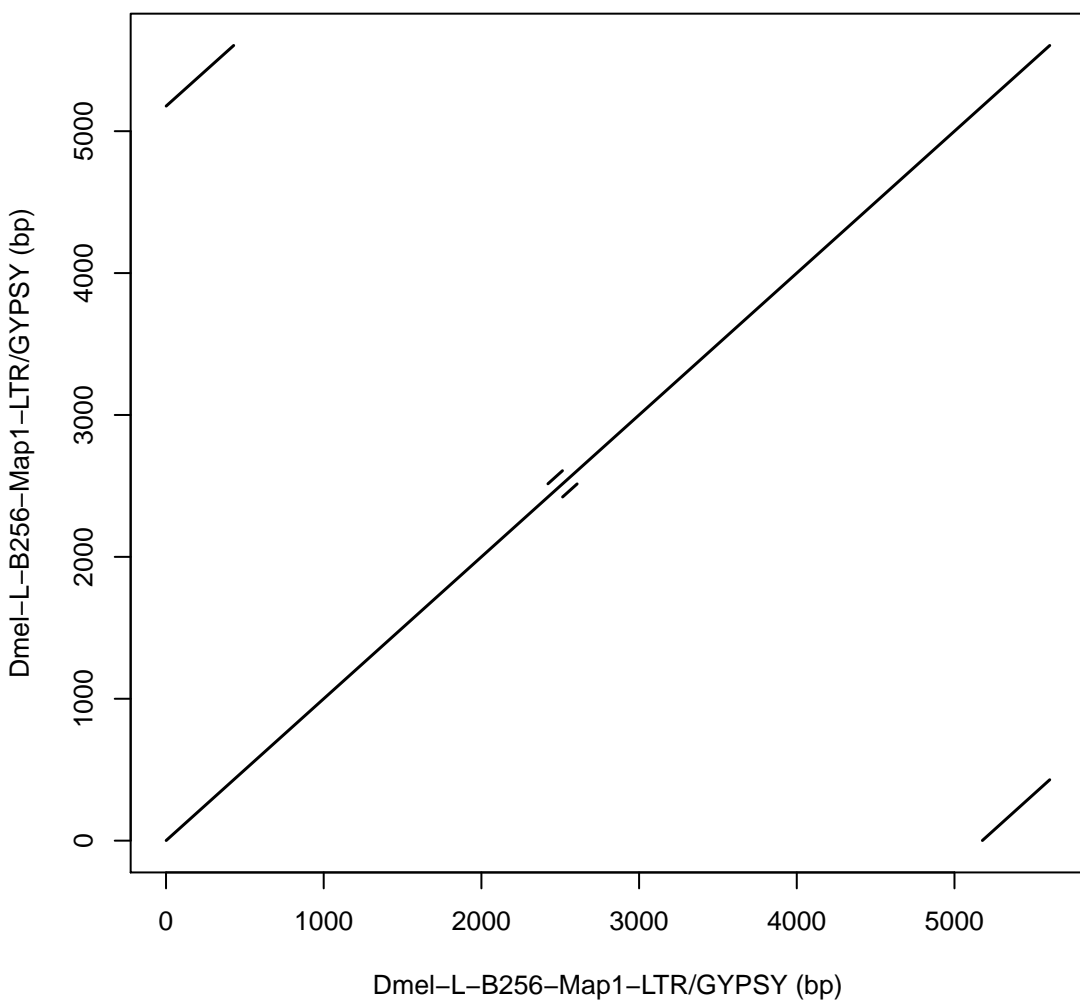
TE: Dmel-L-B256-Map1-LTR/GYPSY
consensus size: 5604bp; fragments: 626; full length: 2 (≥ 5043.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

