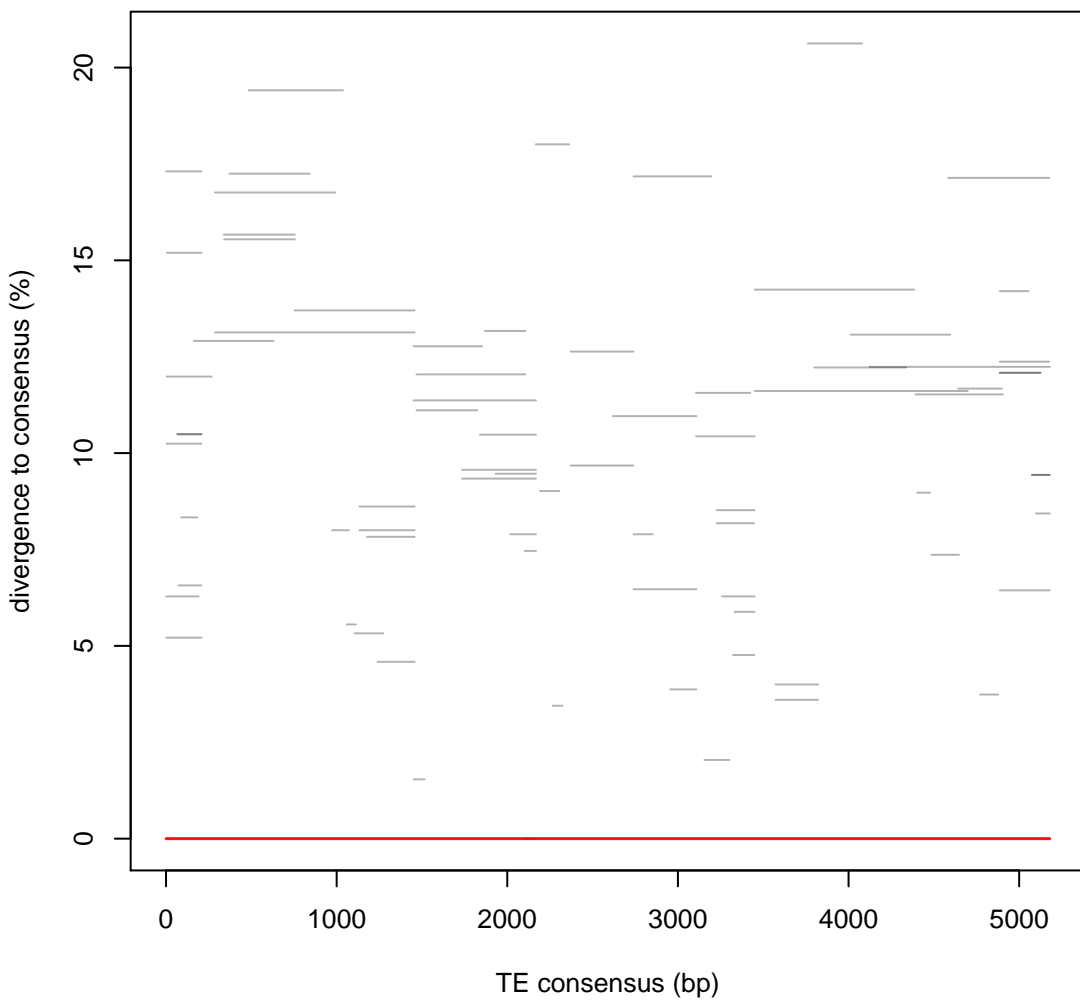
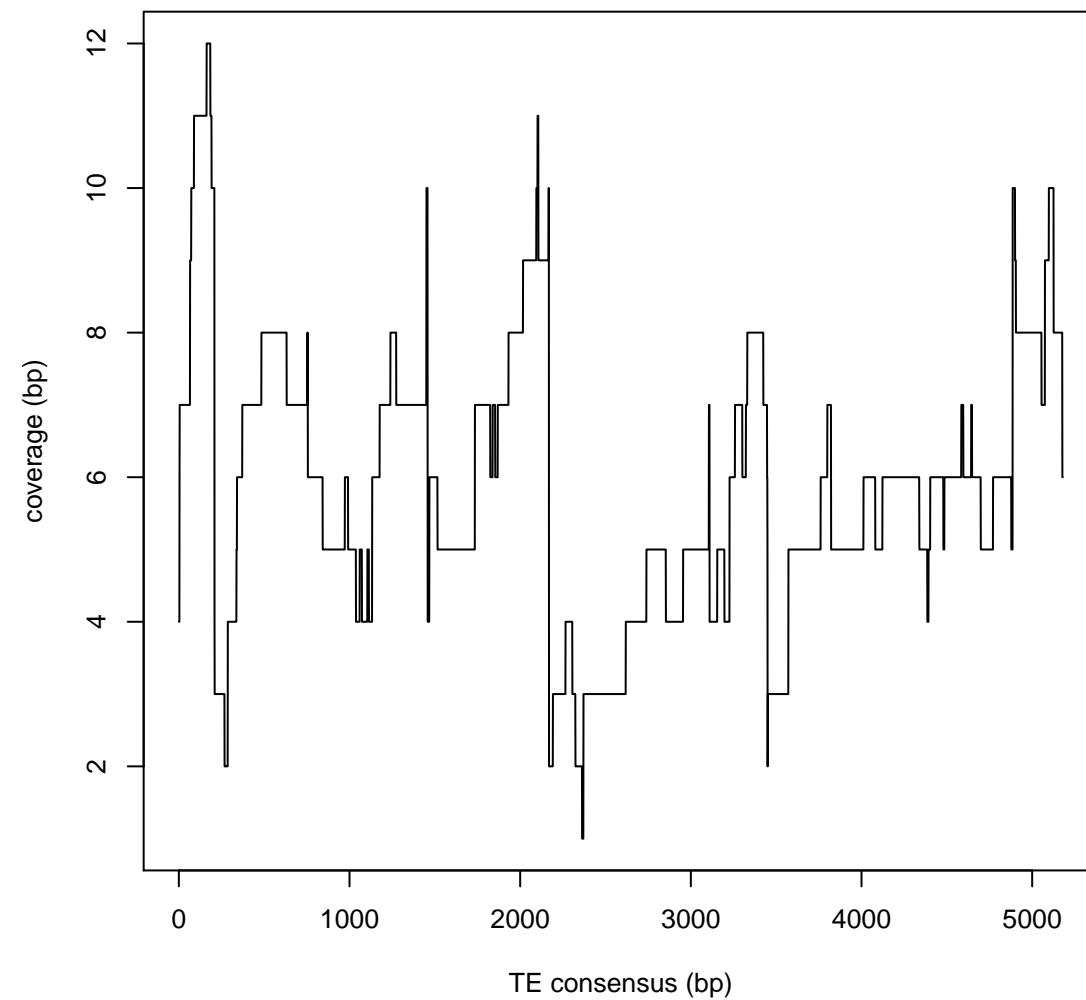


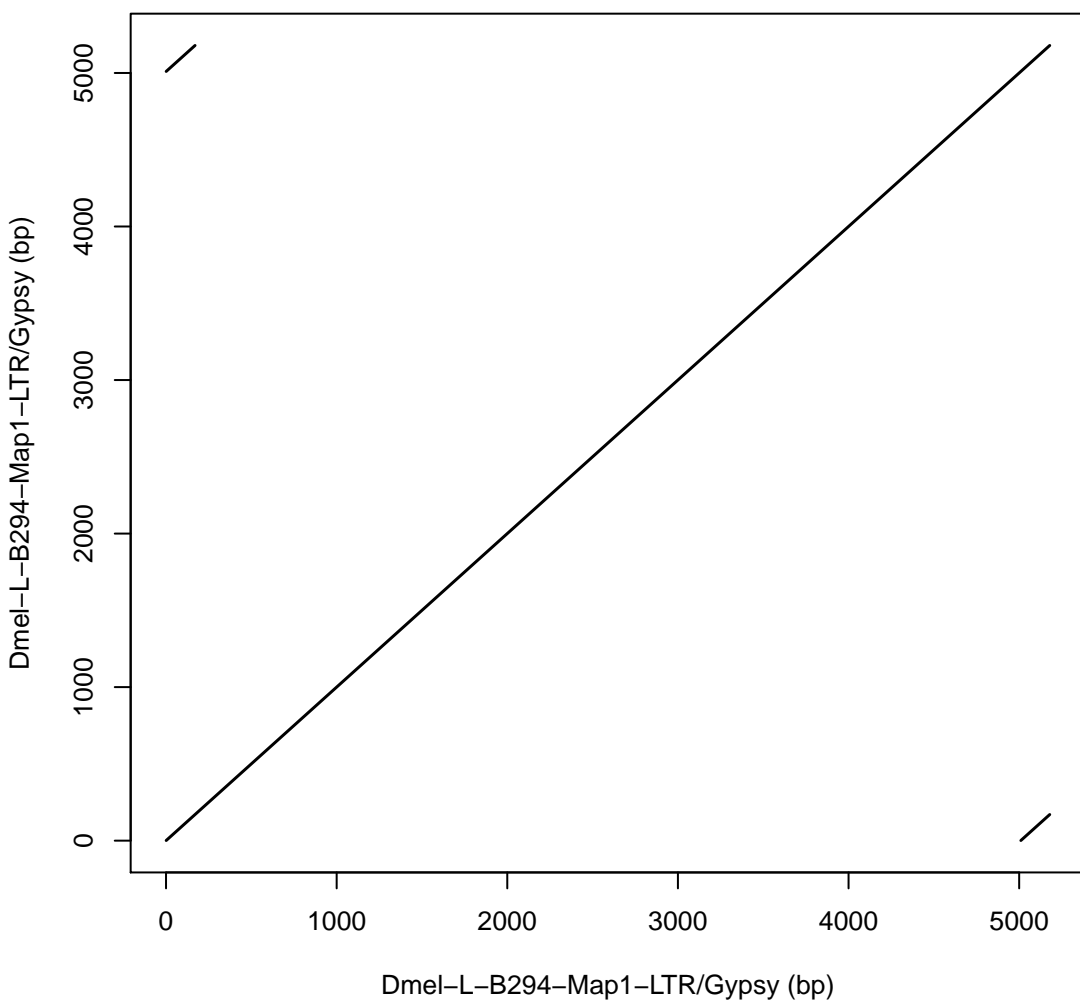
TE: Dmel-L-B294-Map1-LTR/Gypsy
consensus size: 5179bp; fragments: 79; full length: 1 (≥ 4661.1 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

