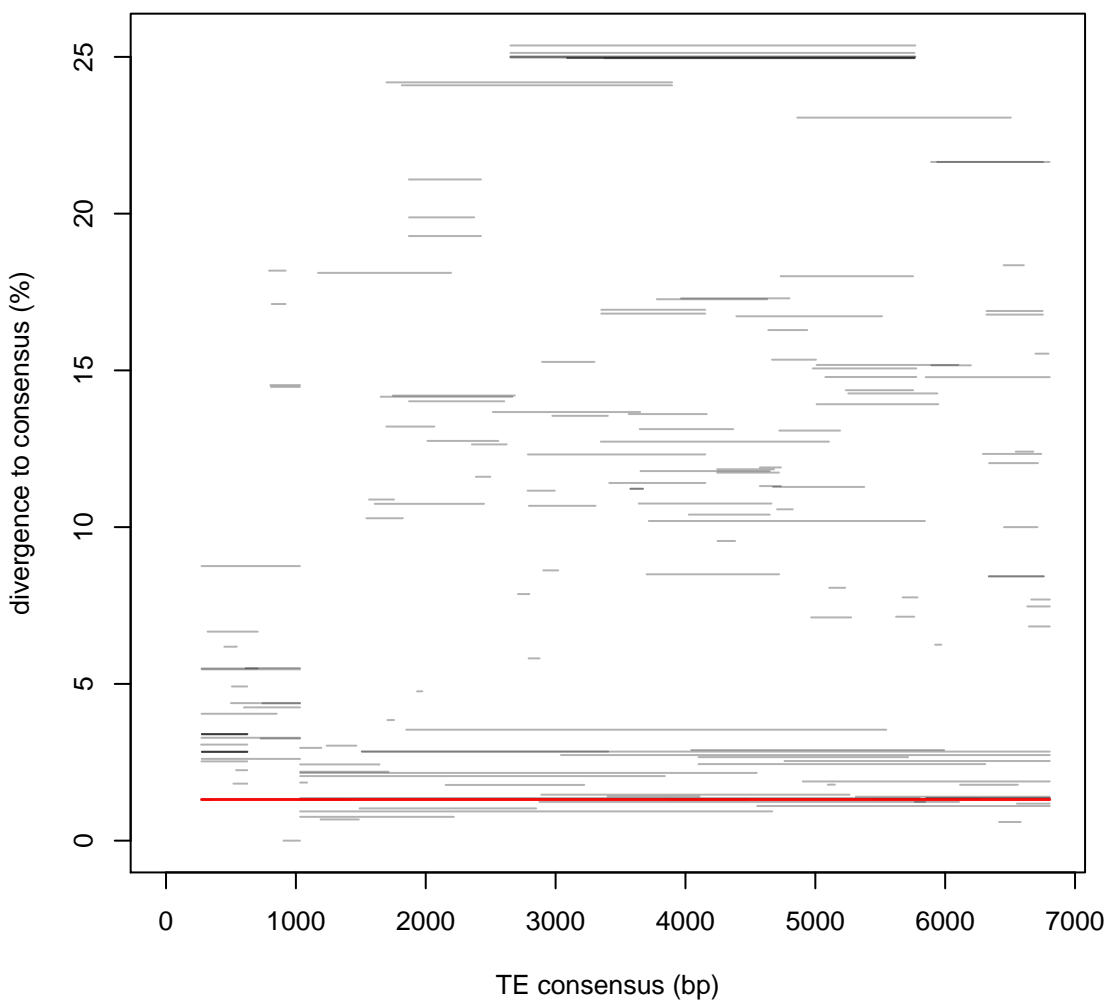
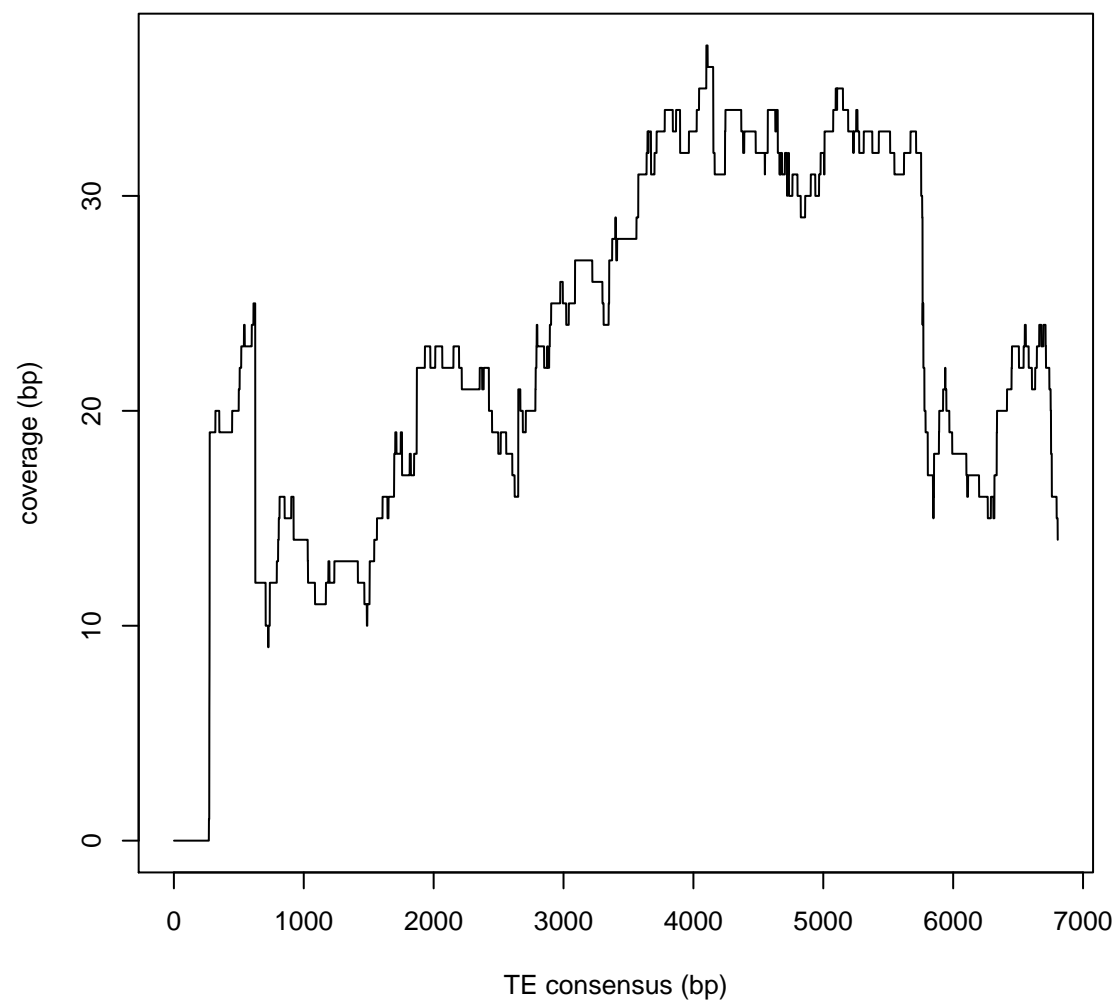


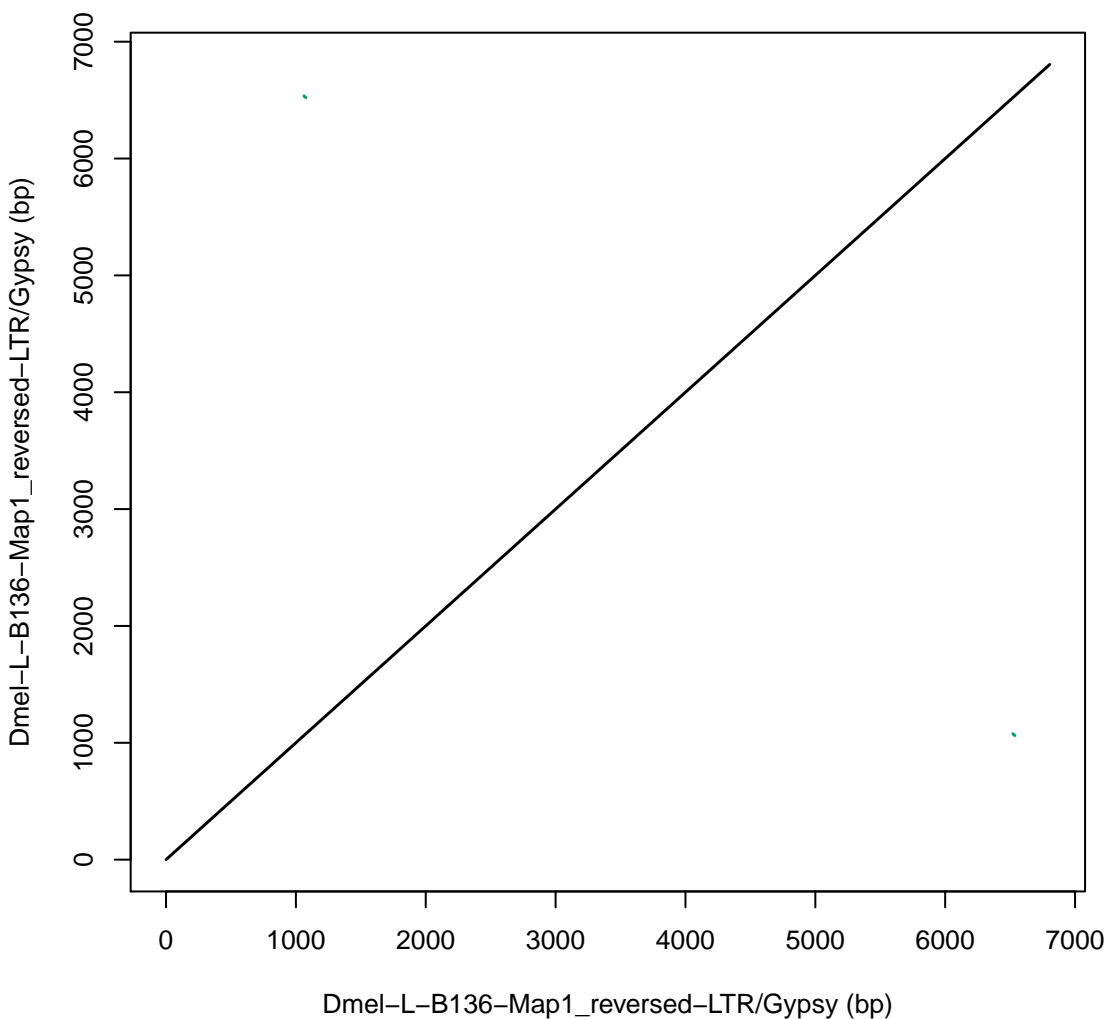
TE: Dmel-L-B136-Map1\_reversed-LTR/Gypsy  
consensus size: 6805bp; fragments: 163; full length: 1 (>=6124.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

