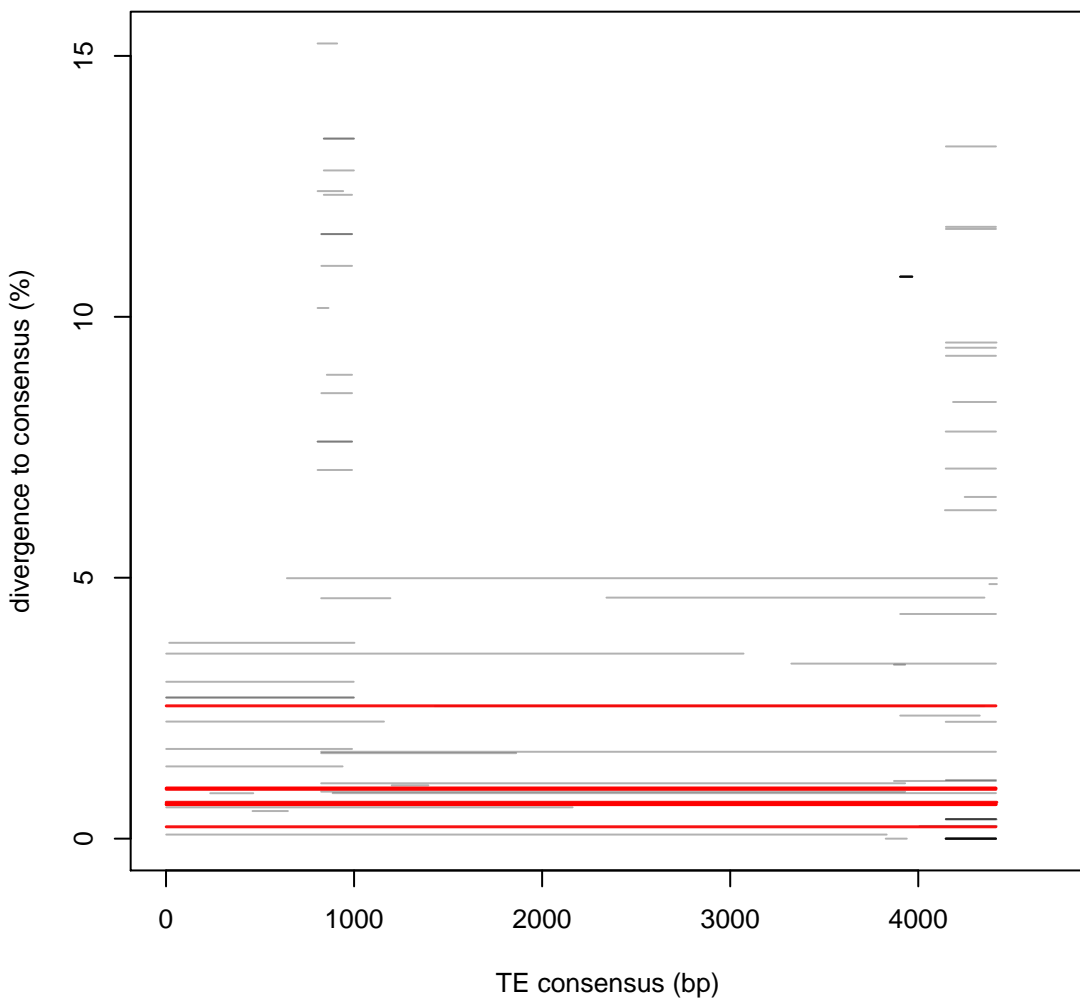
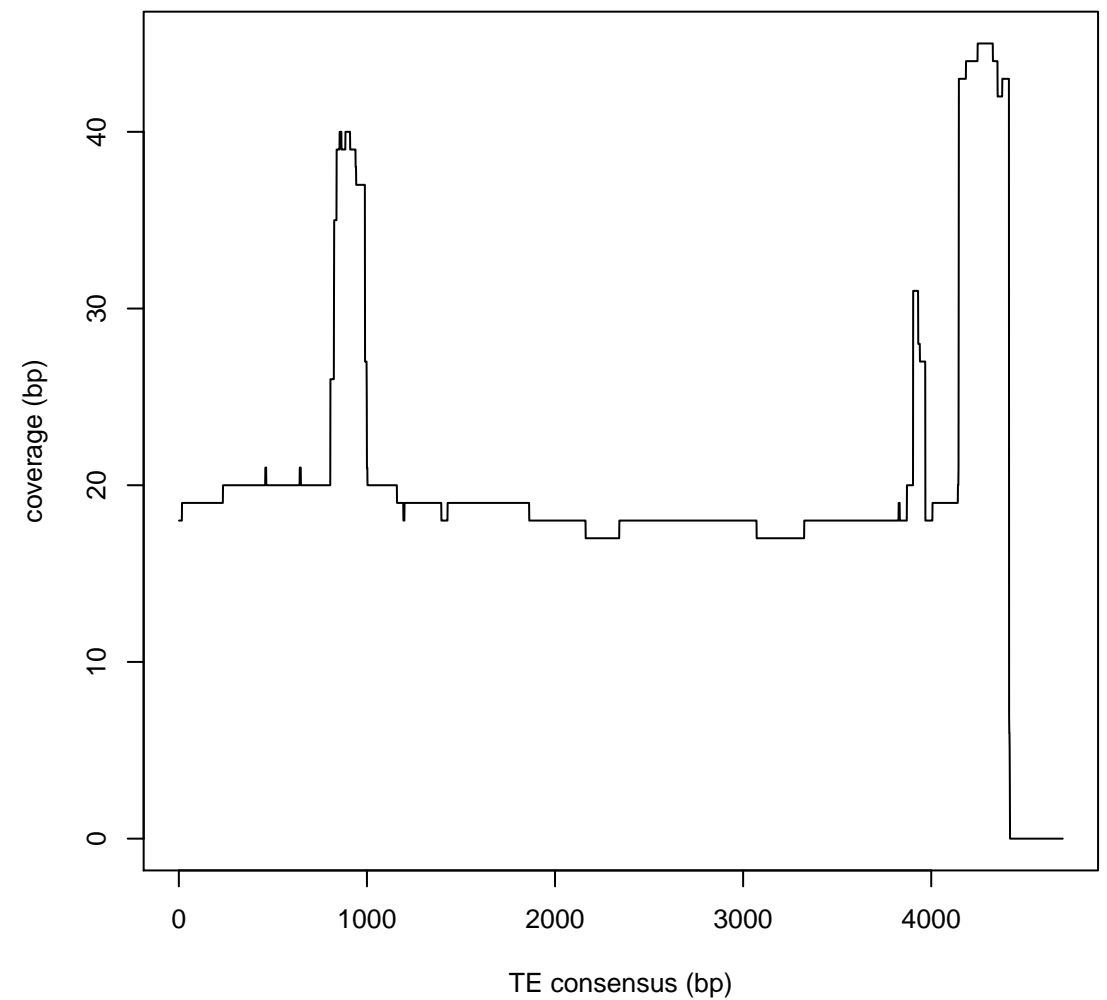


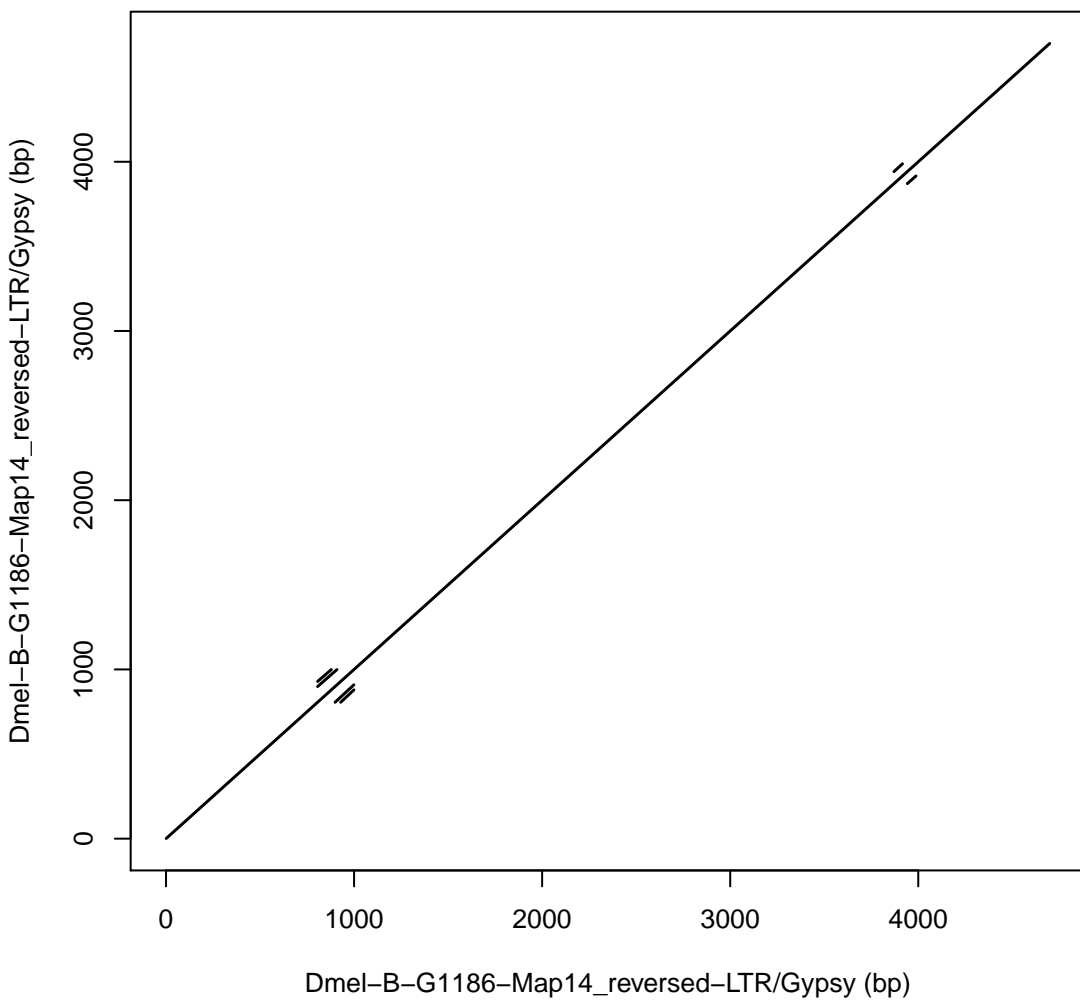
TE: Dmel-B-G1186-Map14_reversed-LTR/Gypsy
consensus size: 4699bp; fragments: 89; full length: 9 (≥ 4229.1 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

