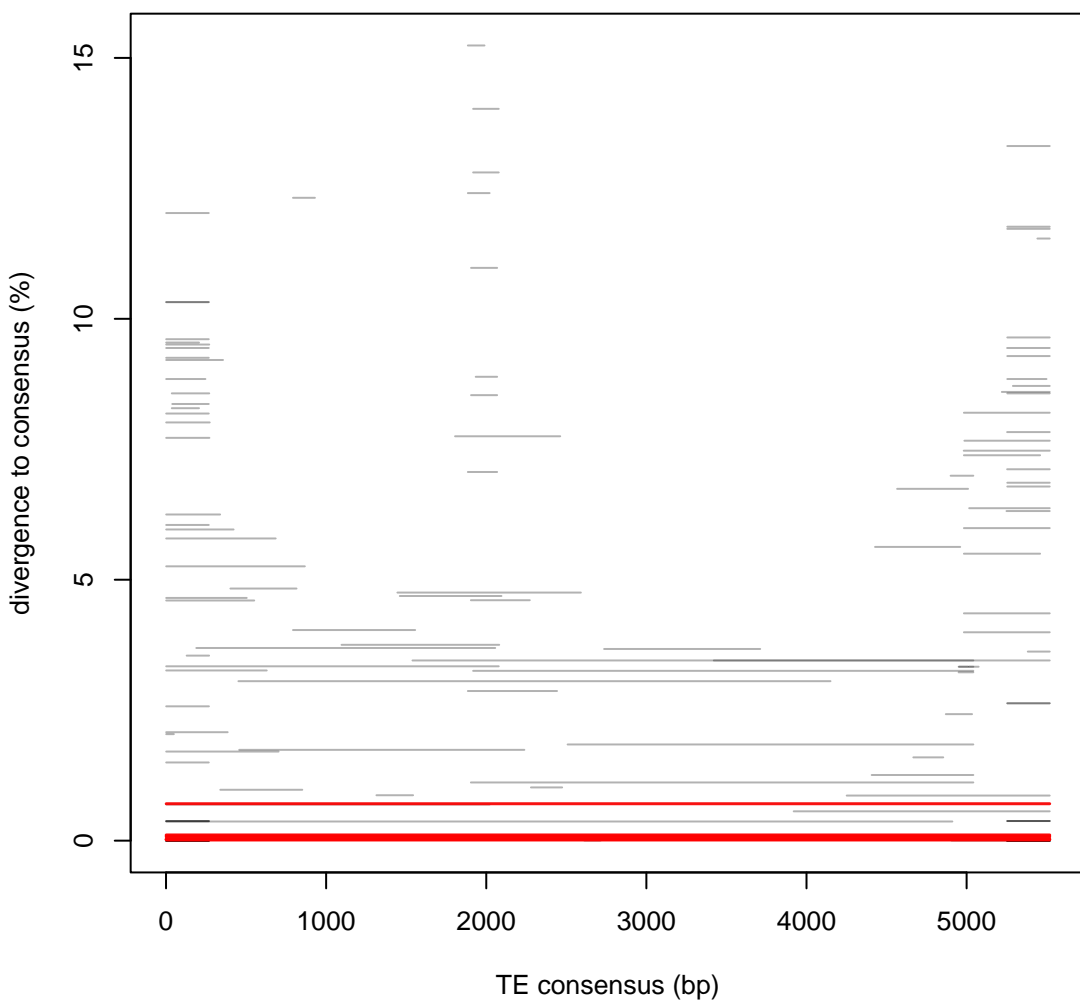
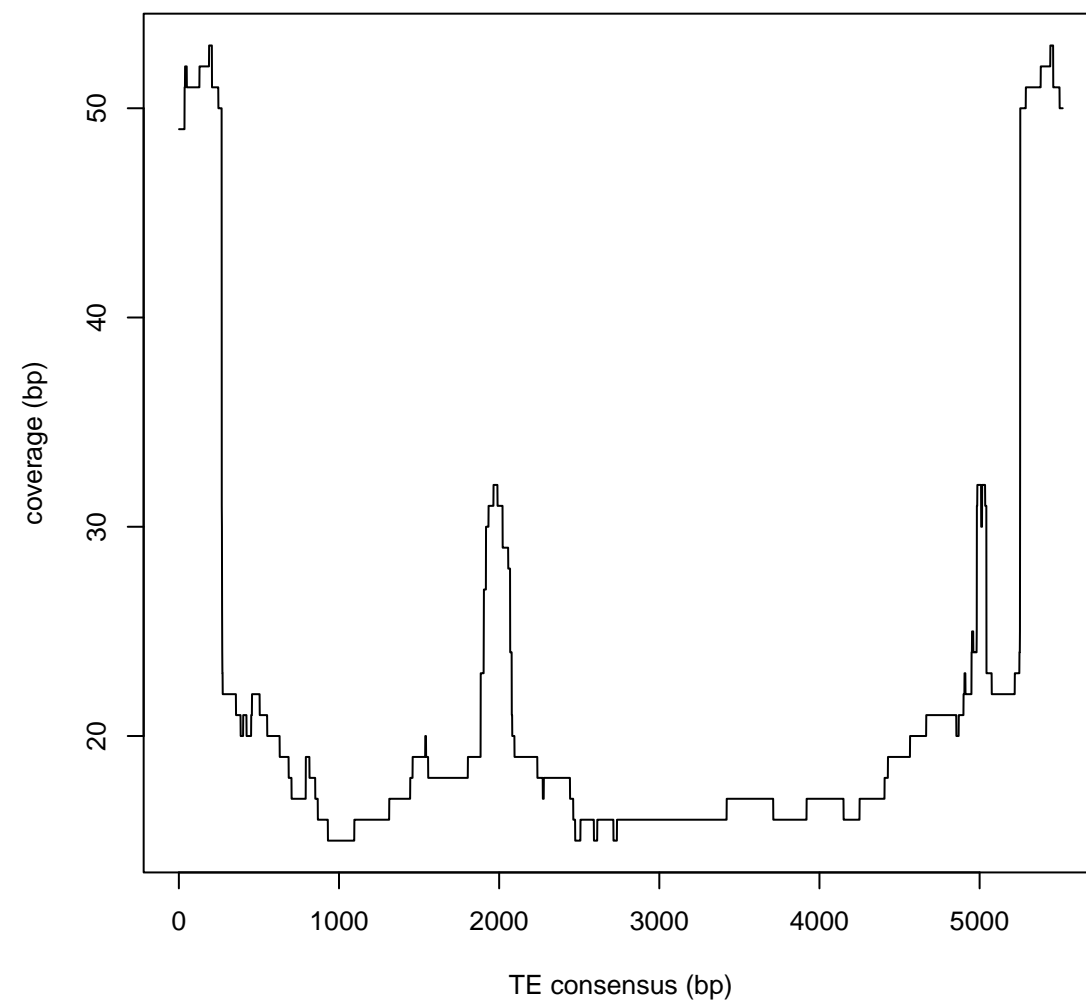


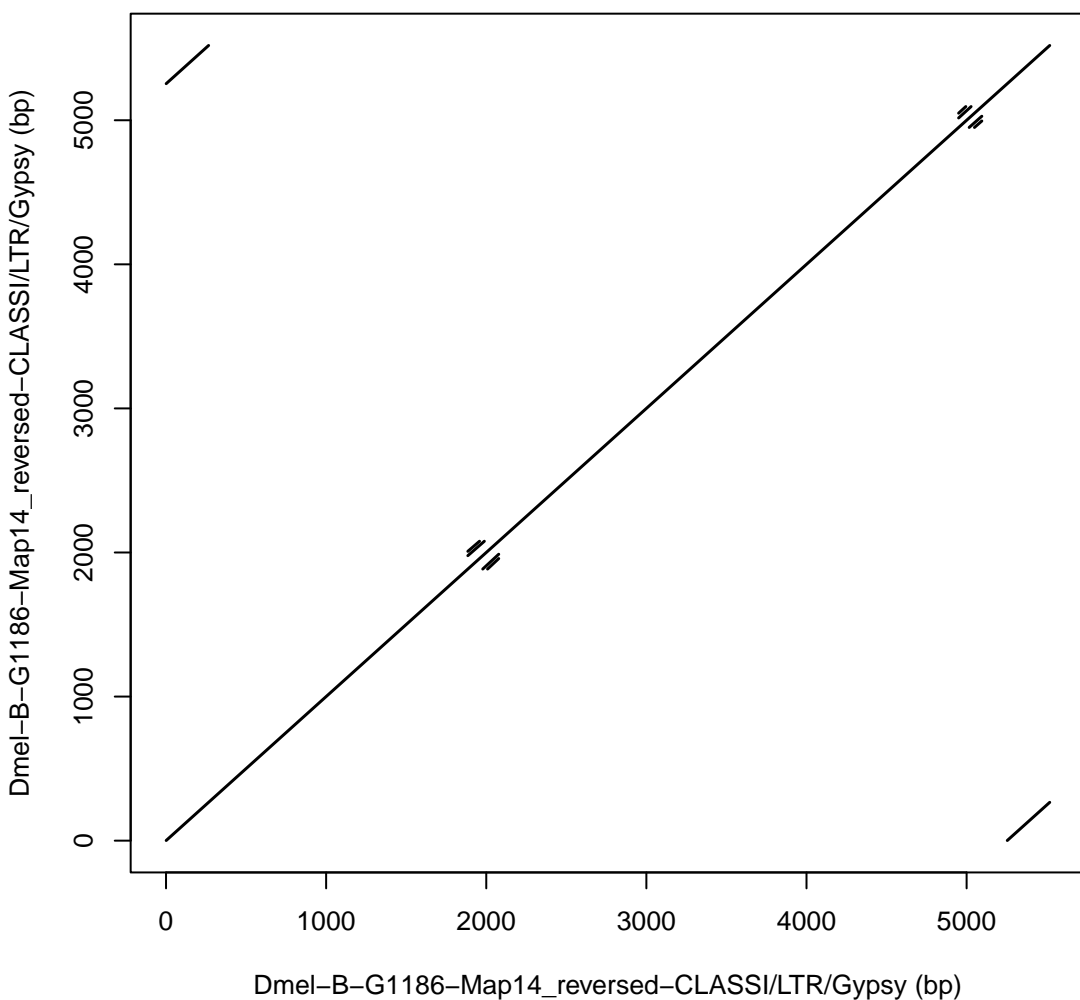
TE: Dmel-B-G1186-Map14_reversed-CLASSI/LTR/Gypsy
consensus size: 5519bp; fragments: 137; full length: 8 (>=4967.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

