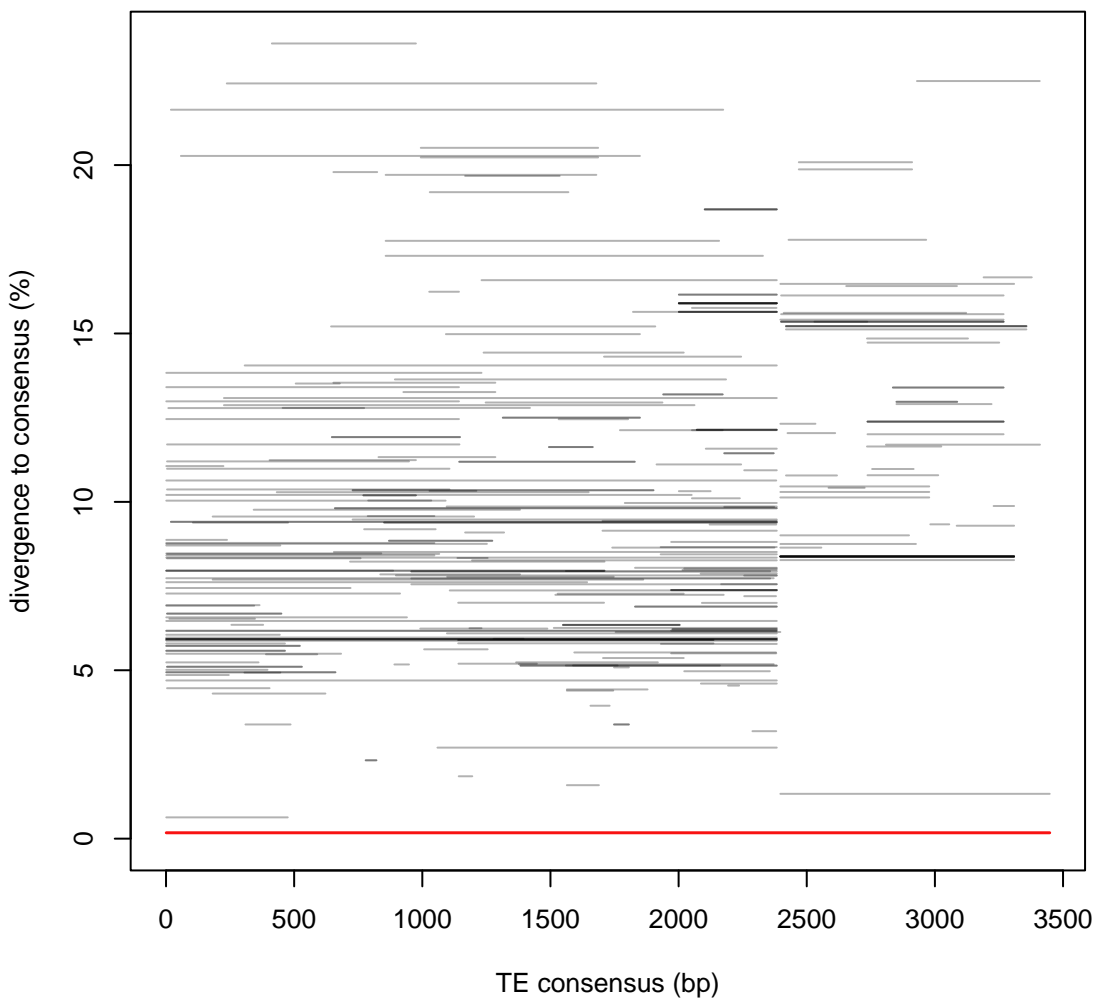
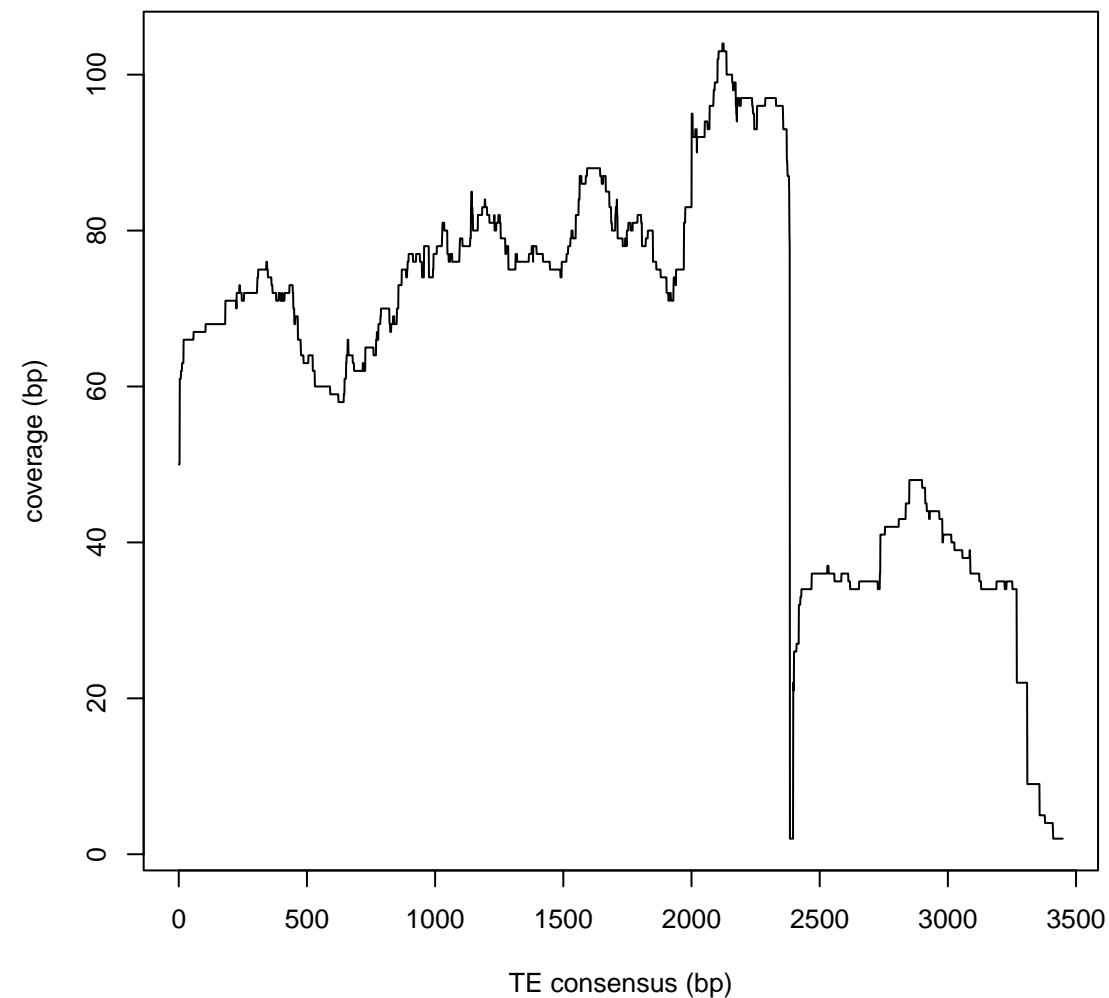


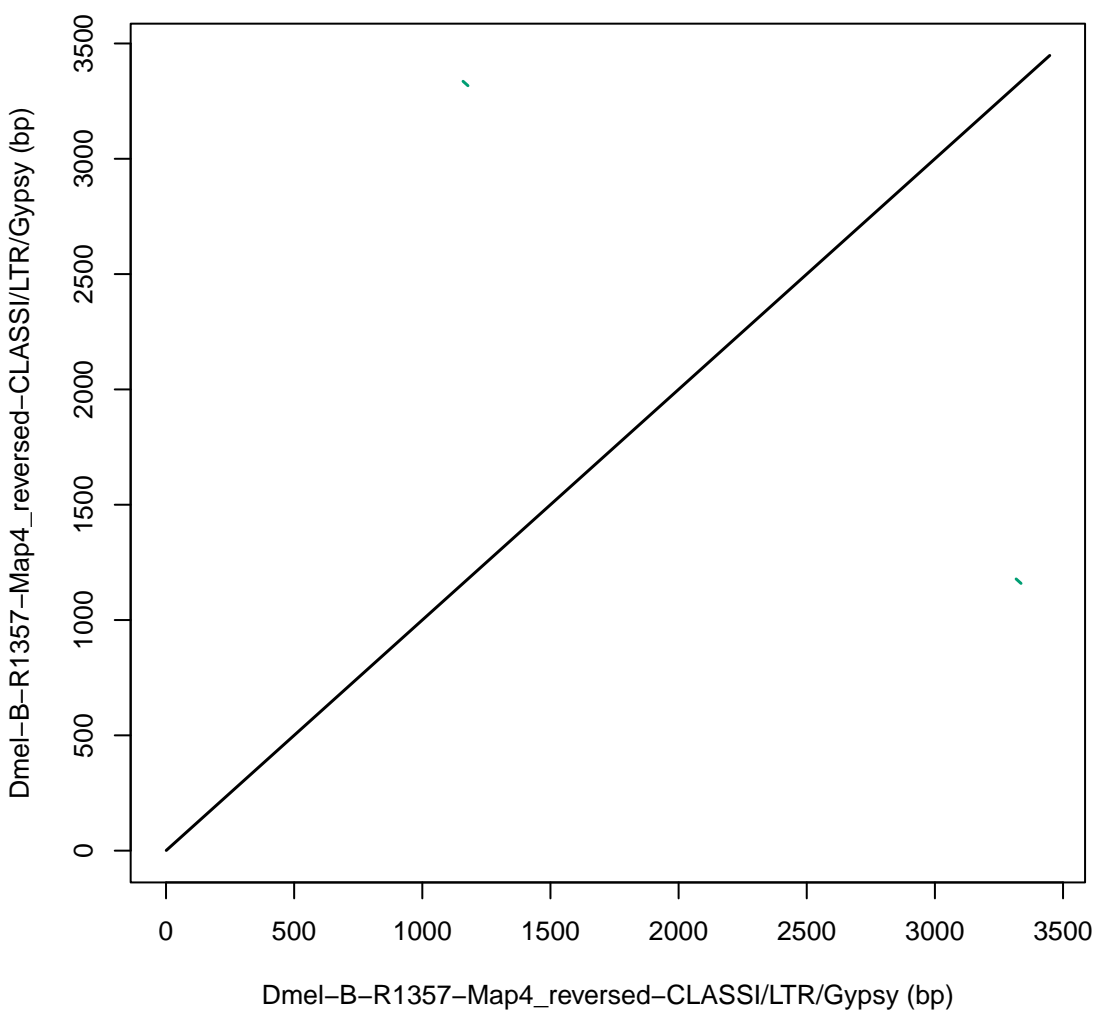
TE: Dmel-B-R1357-Map4_reversed-CLASSI/LTR/Gypsy
consensus size: 3448bp; fragments: 299; full length: 1 (>=3103.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

