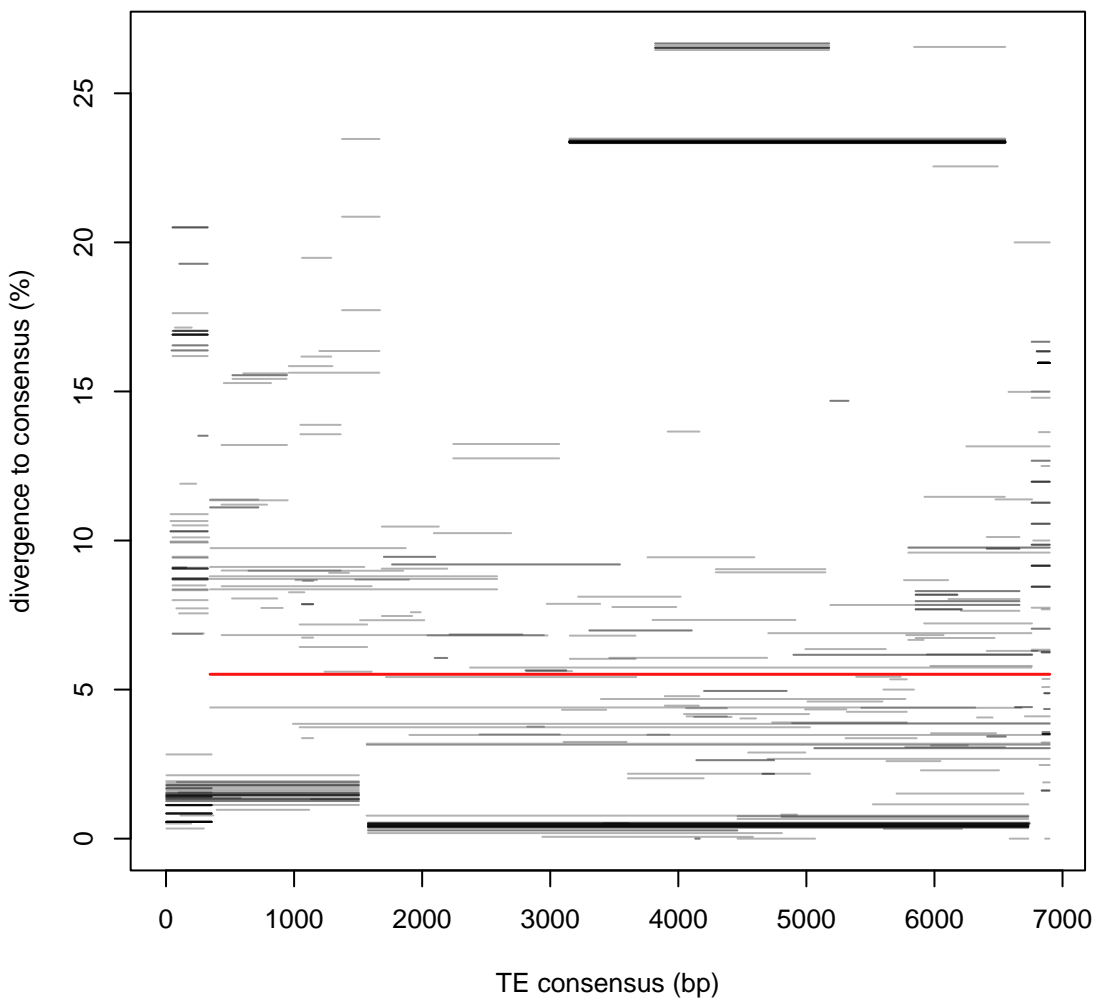
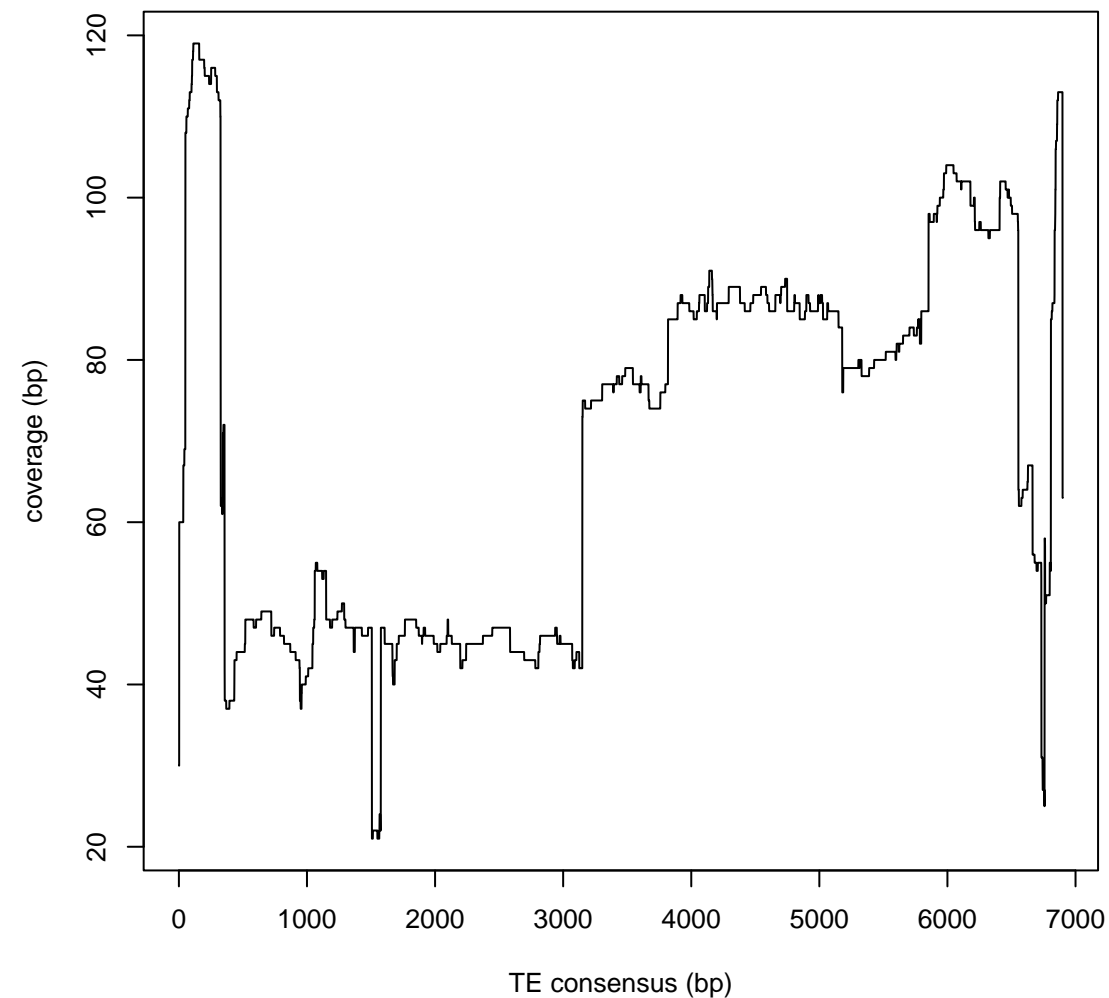


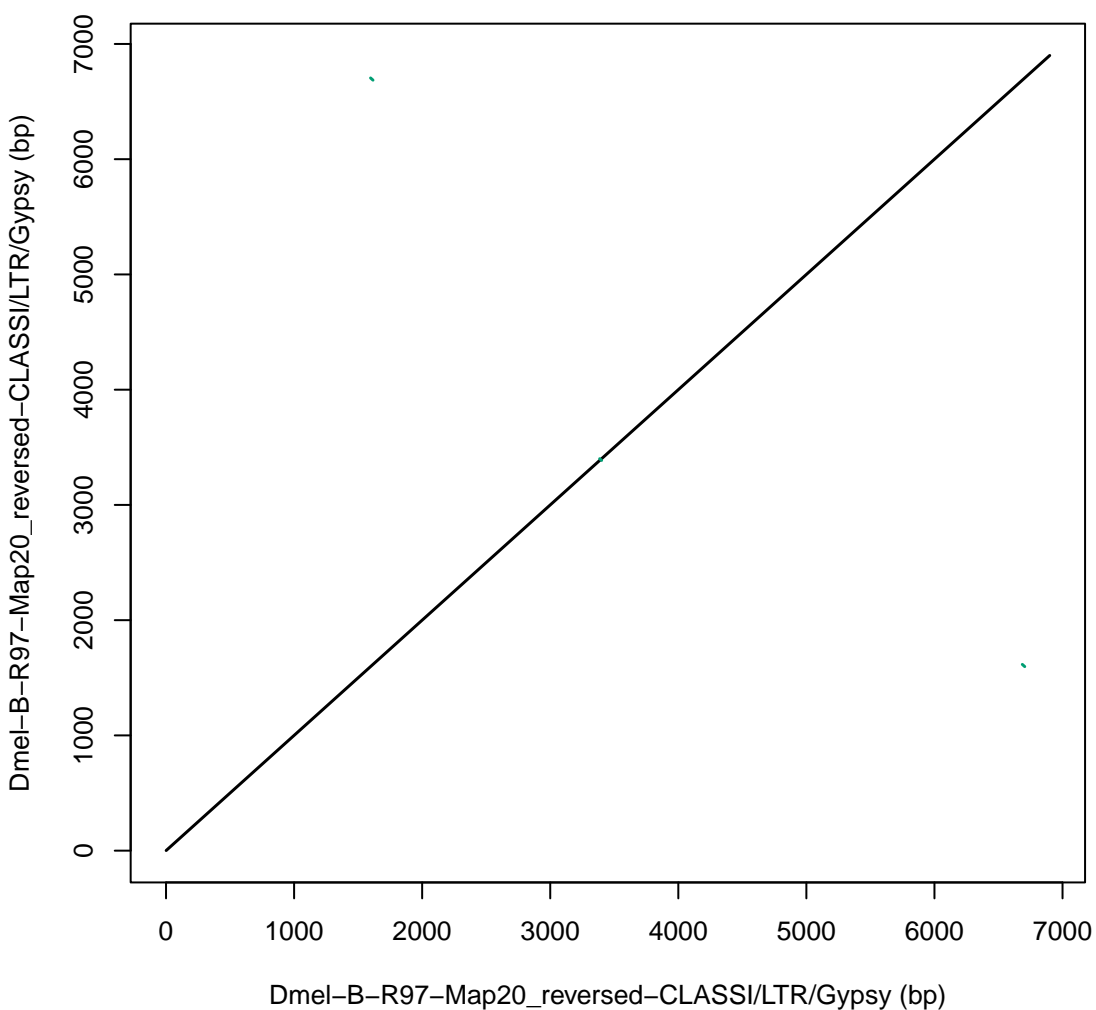
TE: Dmel-B-R97-Map20_reversed-CLASSI/LTR/Gypsy
consensus size: 6900bp; fragments: 471; full length: 1 (>=6210bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

