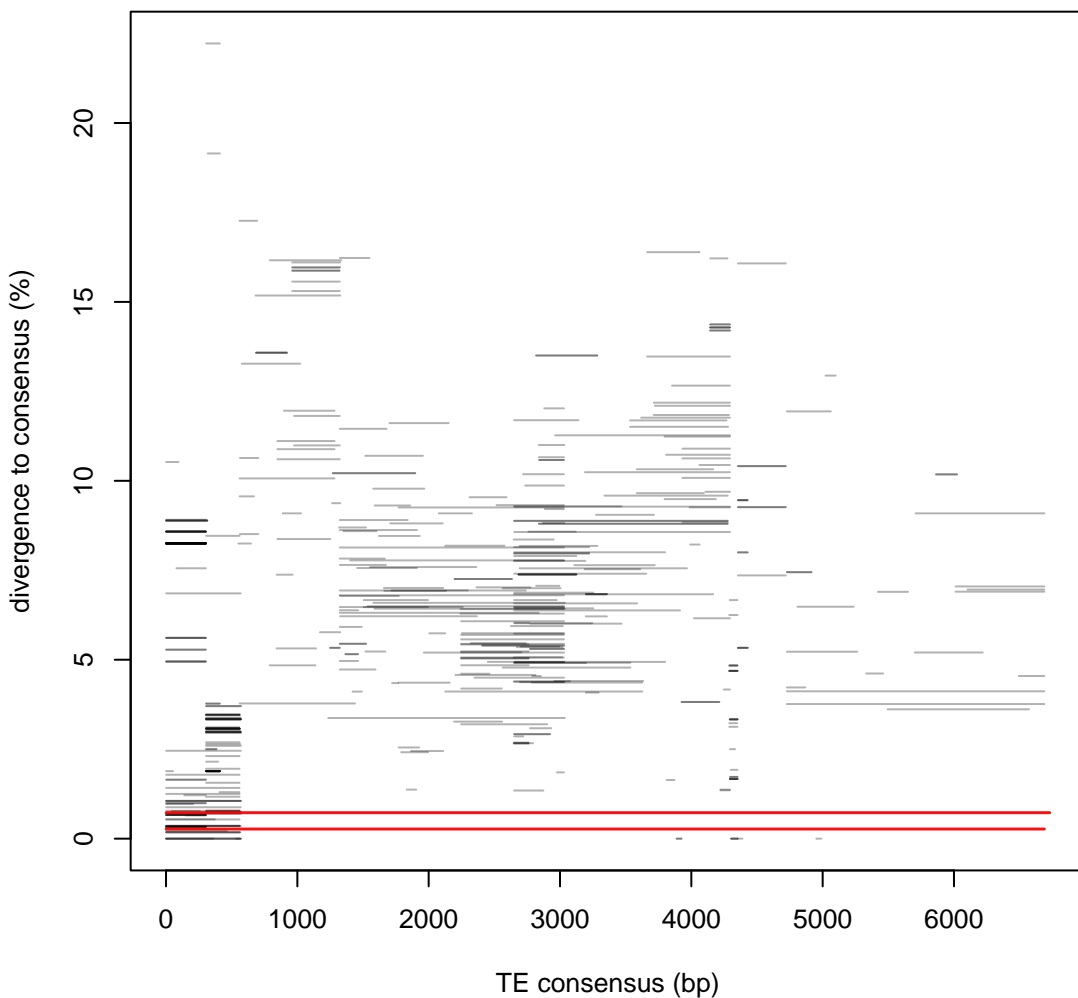
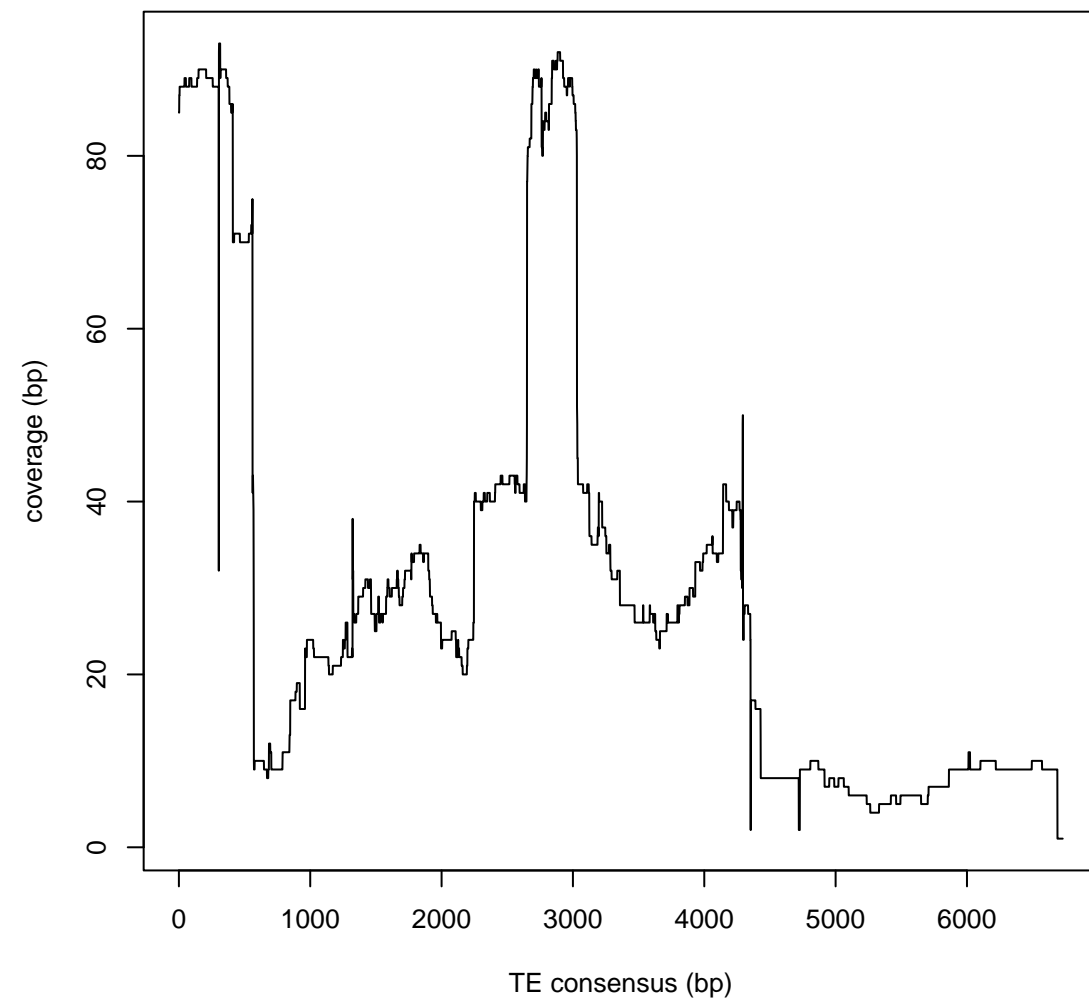


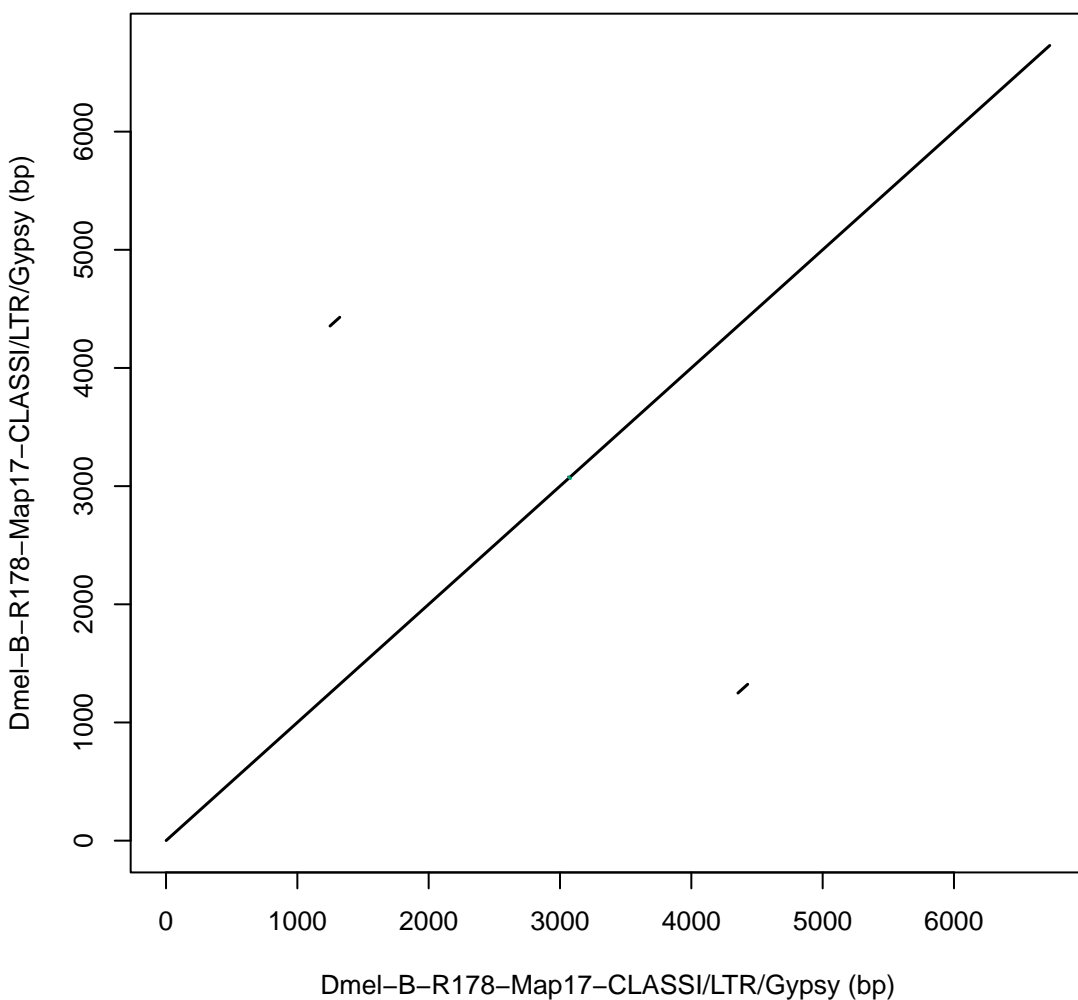
TE: Dmel-B-R178-Map17-CLASSI/LTR/Gypsy
consensus size: 6729bp; fragments: 466; full length: 2 (>=6056.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

