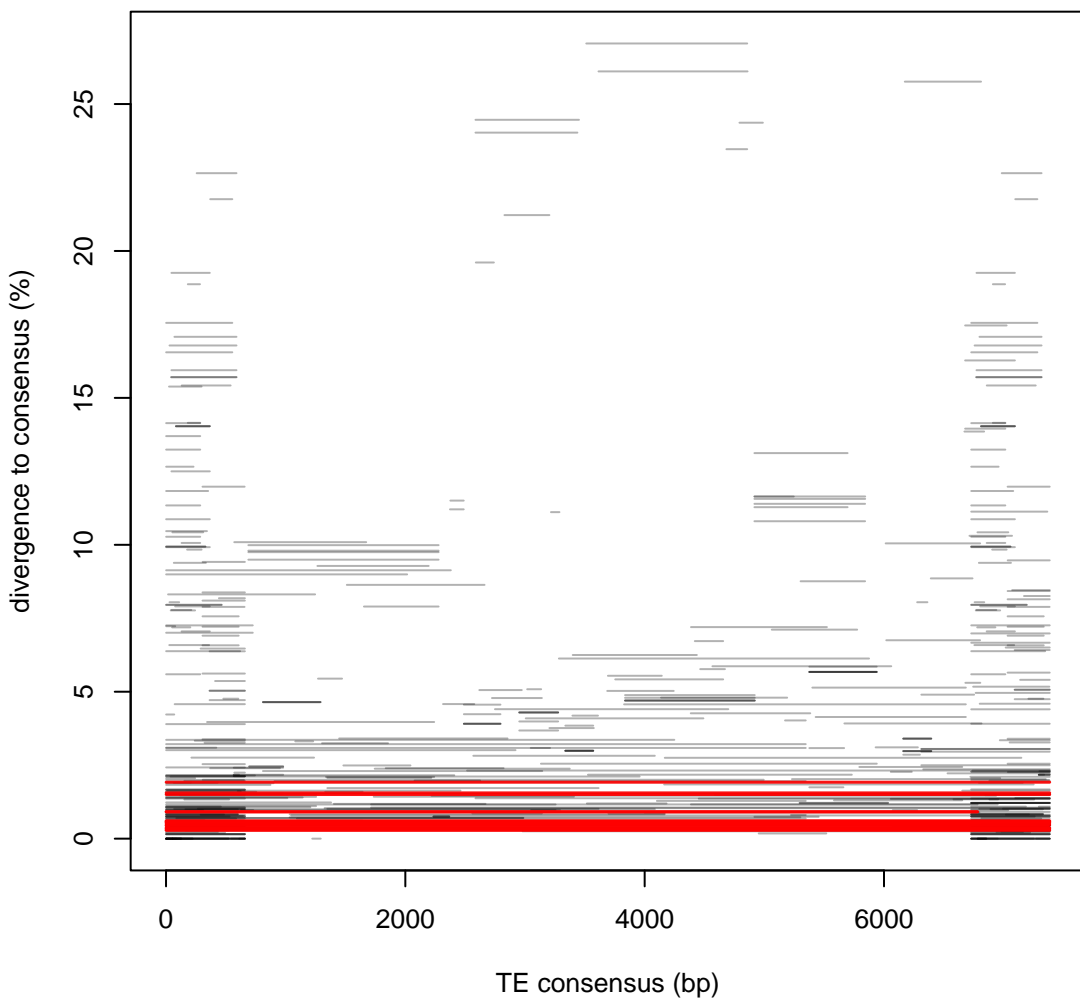
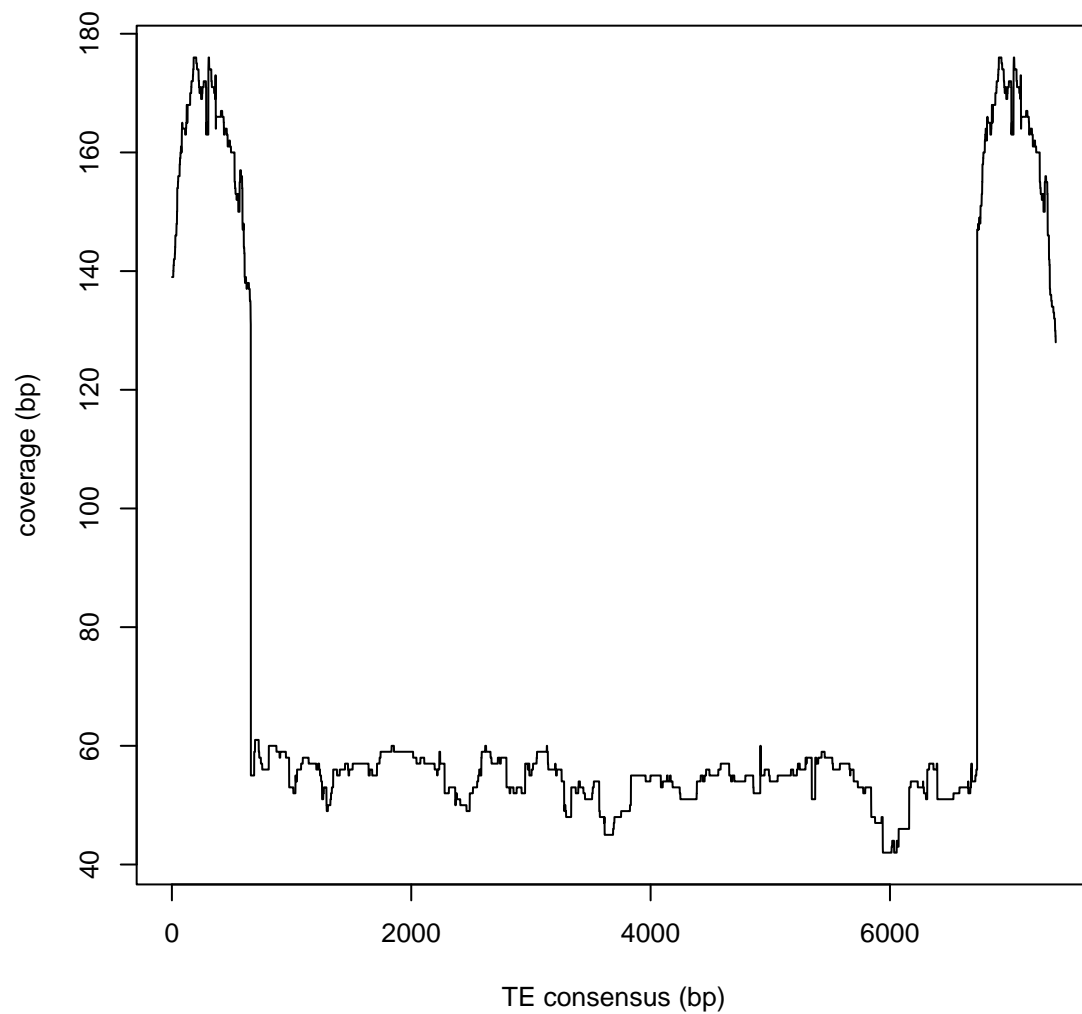


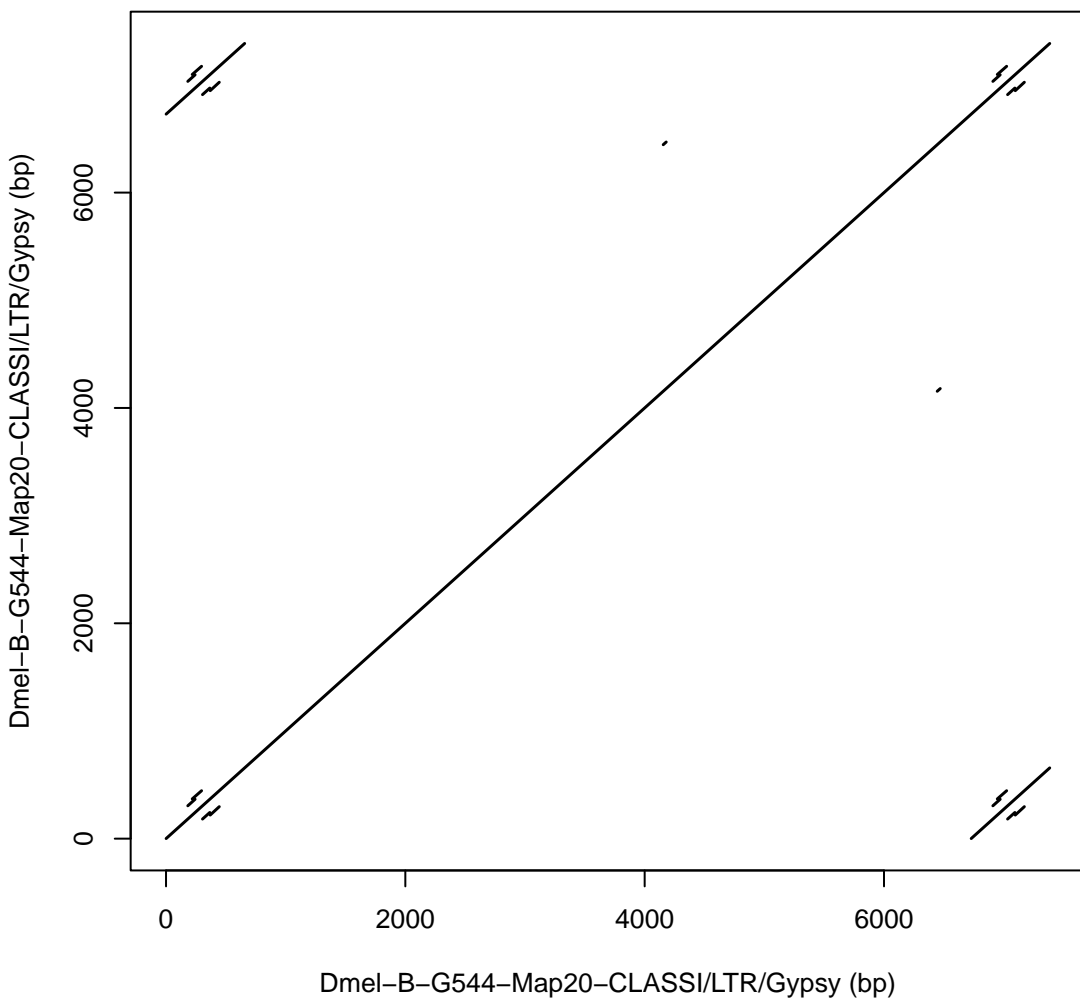
TE: Dmel-B-G544-Map20-CLASSI/LTR/Gypsy
consensus size: 7385bp; fragments: 578; full length: 16 (>=6646.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

