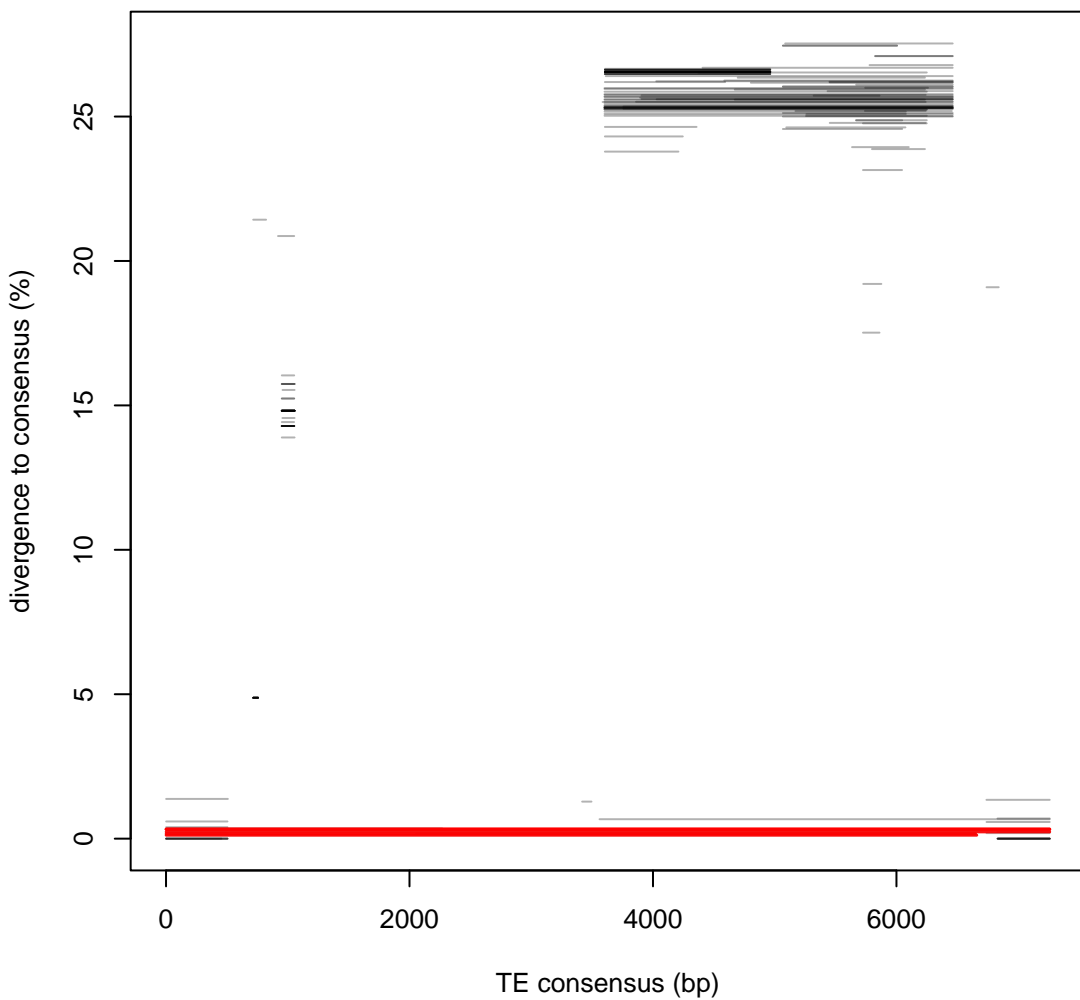
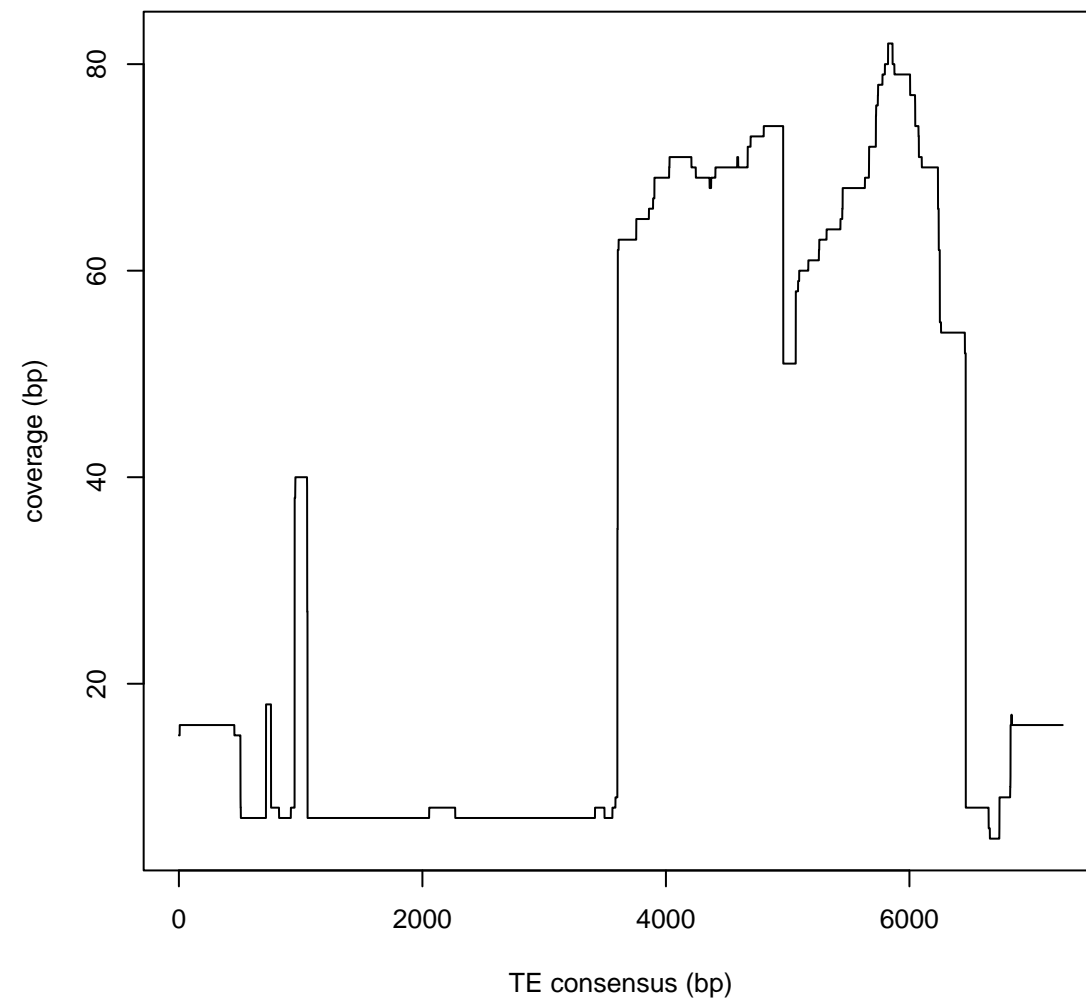


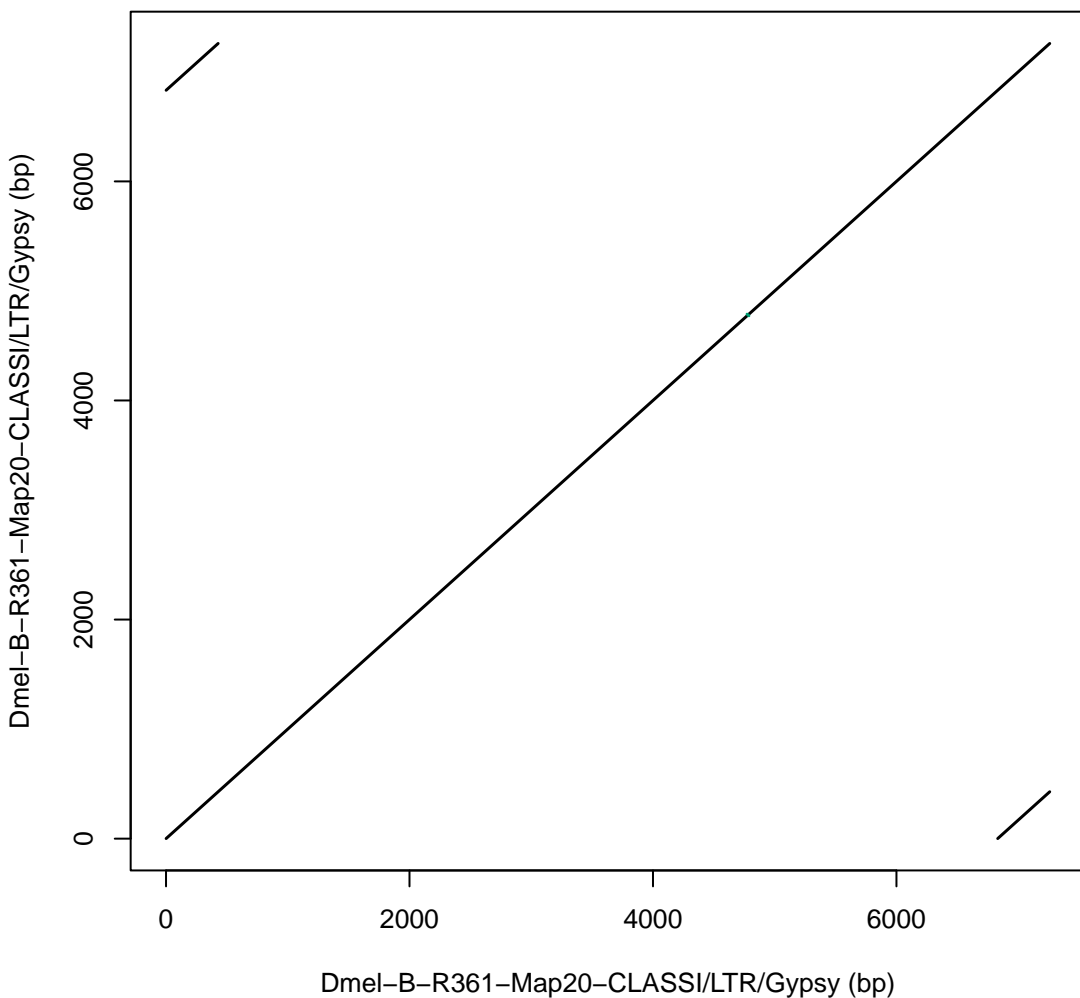
TE: Dmel-B-R361-Map20-CLASSI/LTR/Gypsy
consensus size: 7259bp; fragments: 176; full length: 6 (>=6533.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

