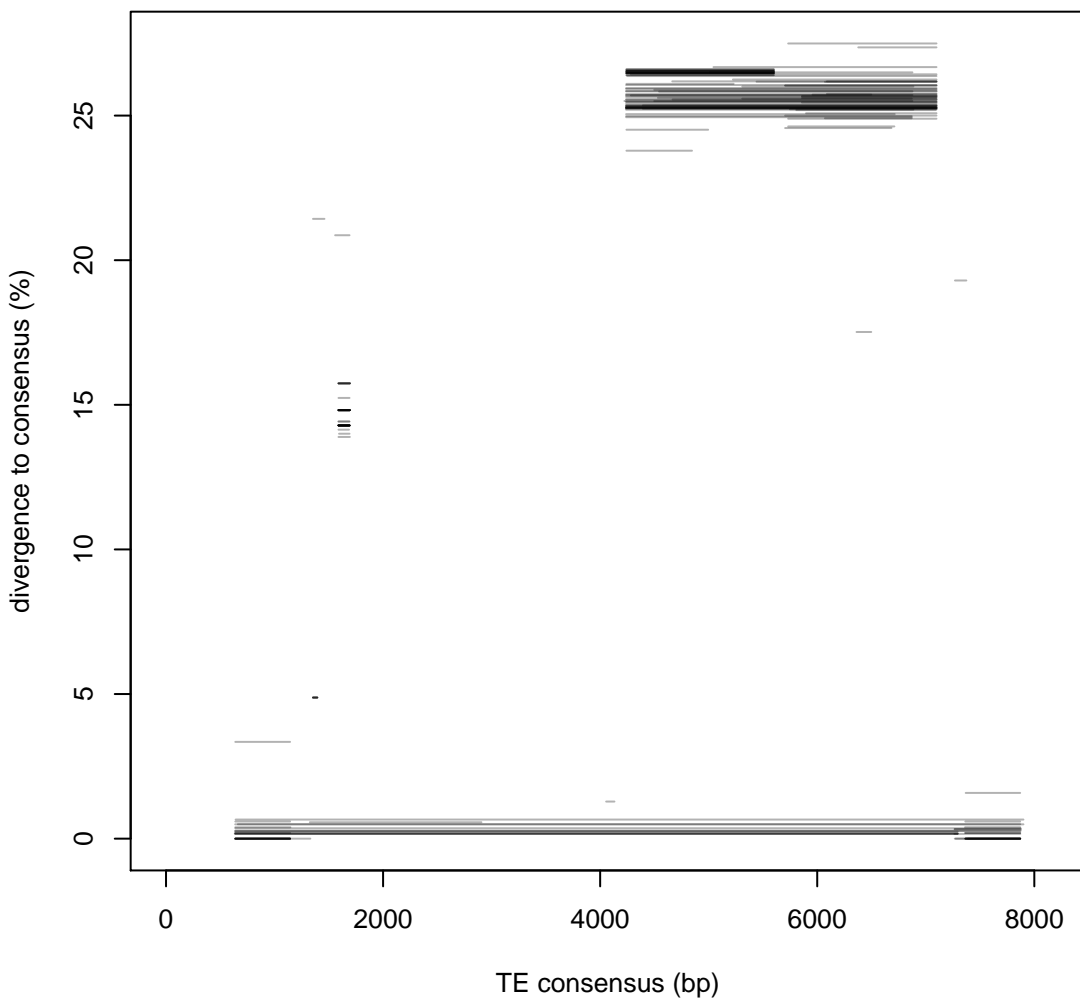
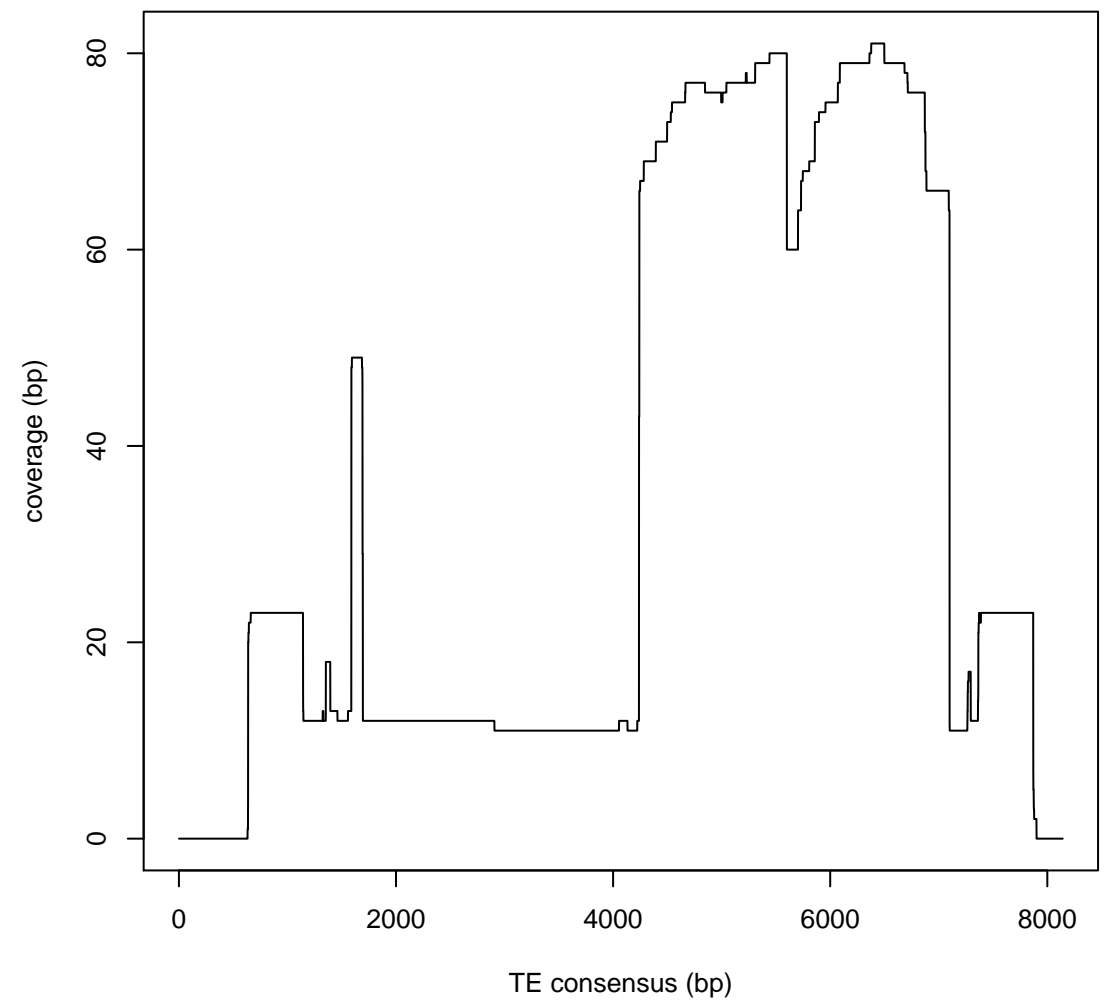


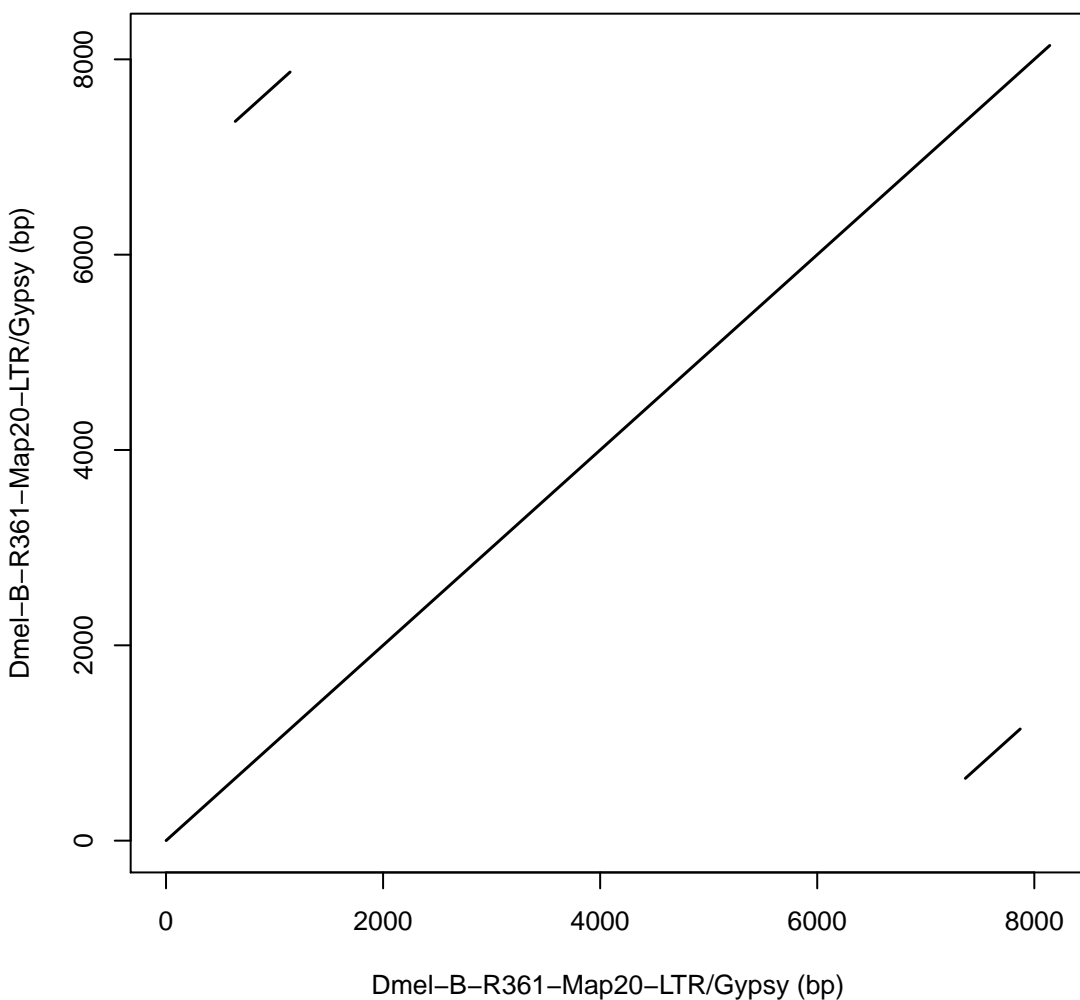
TE: Dmel-B-R361-Map20-LTR/Gypsy
consensus size: 8142bp; fragments: 179; full length: 0 (>=7327.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

