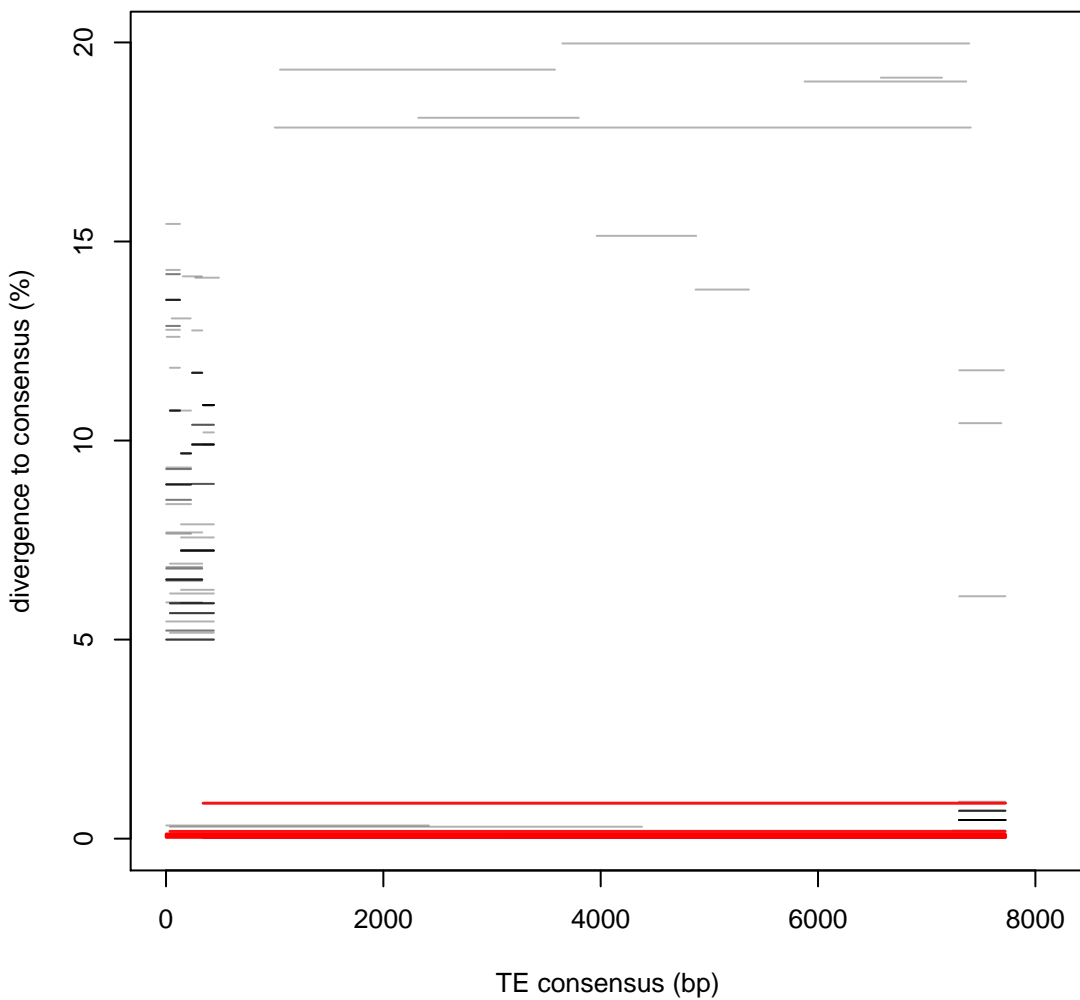
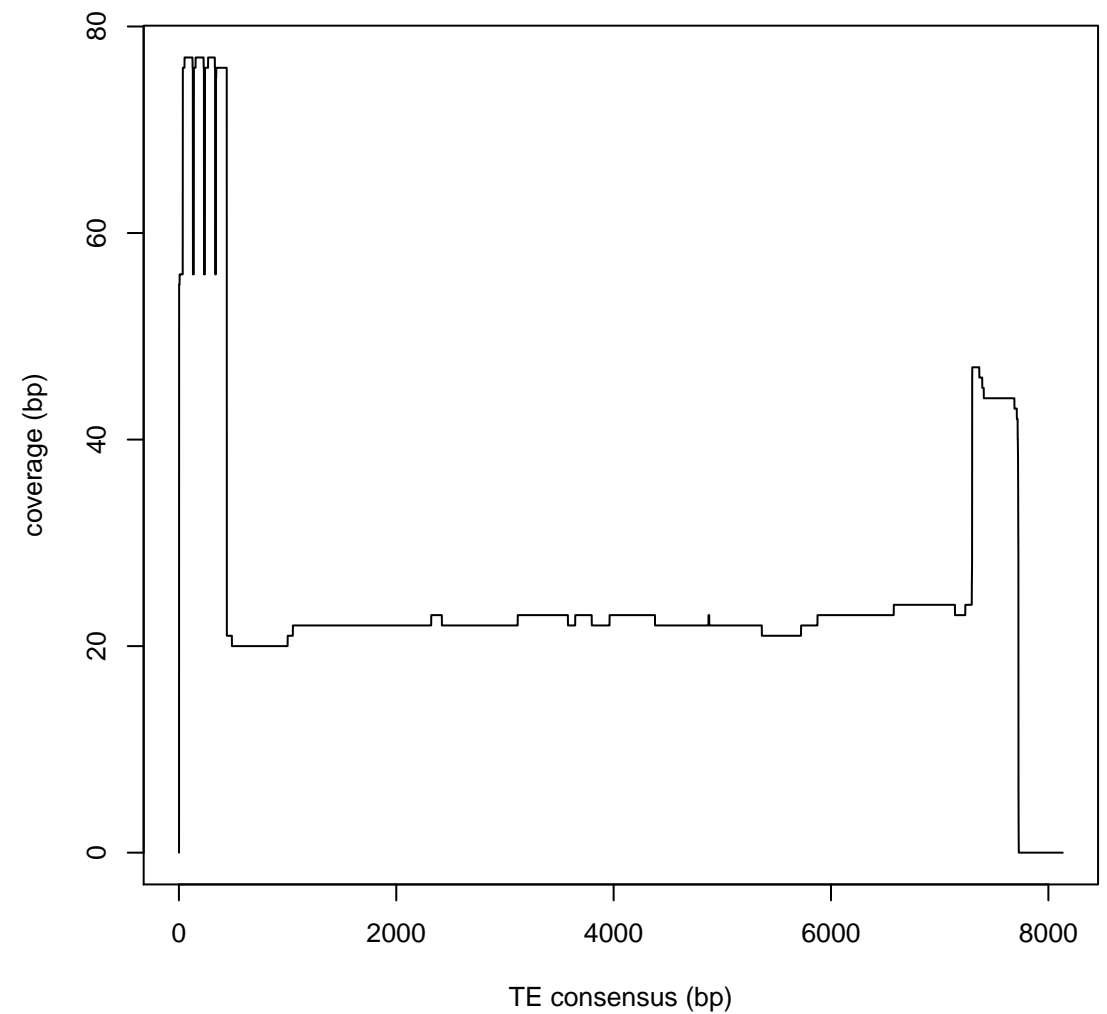


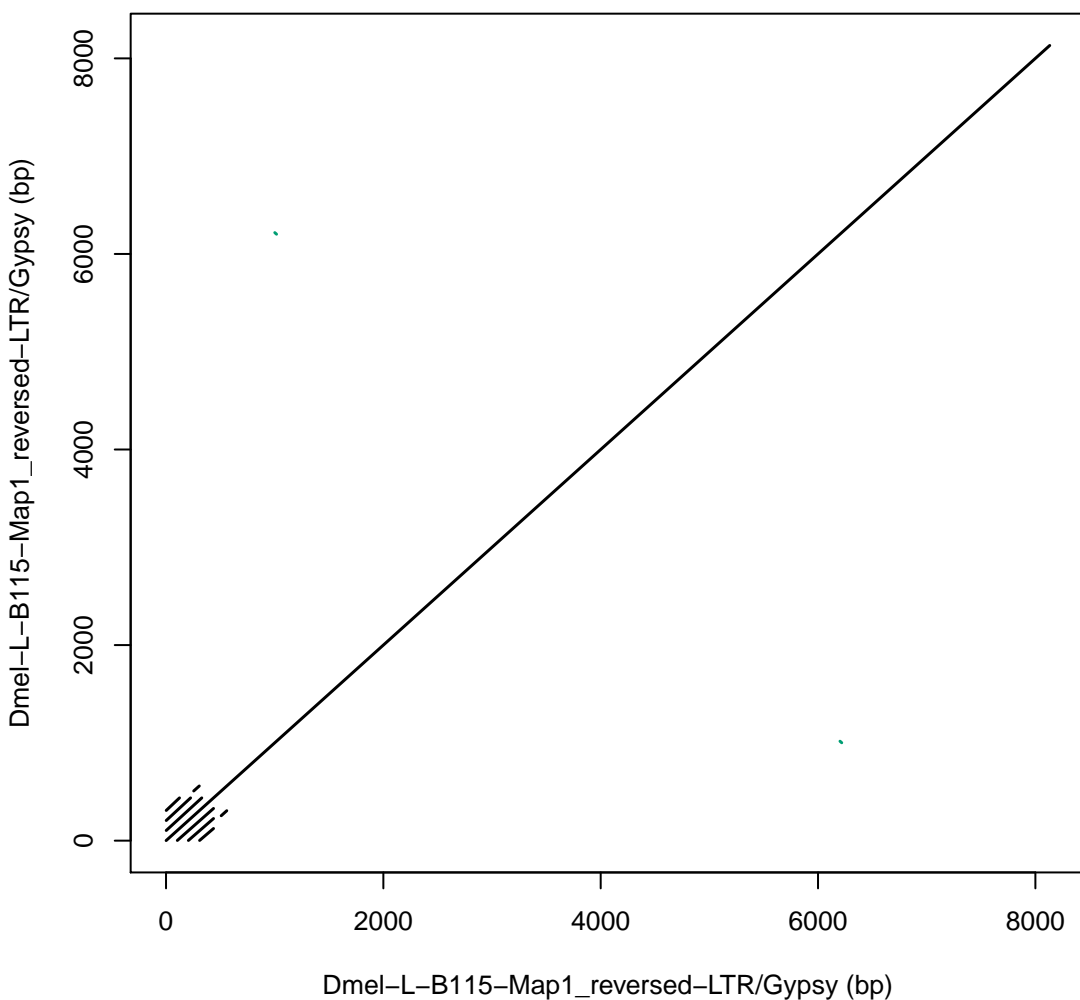
TE: Dmel-L-B115-Map1_reversed-LTR/Gypsy
consensus size: 8132bp; fragments: 173; full length: 18 (>=7318.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

