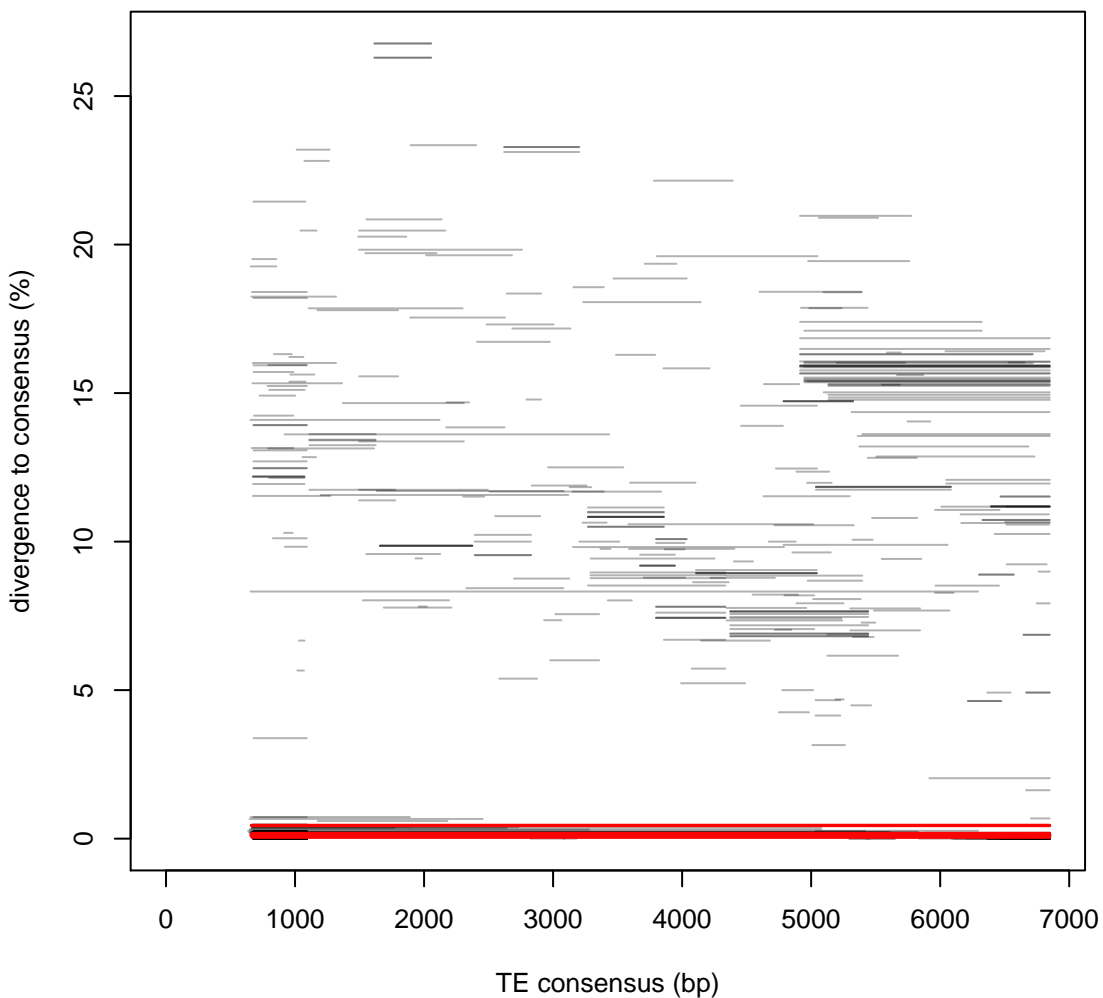
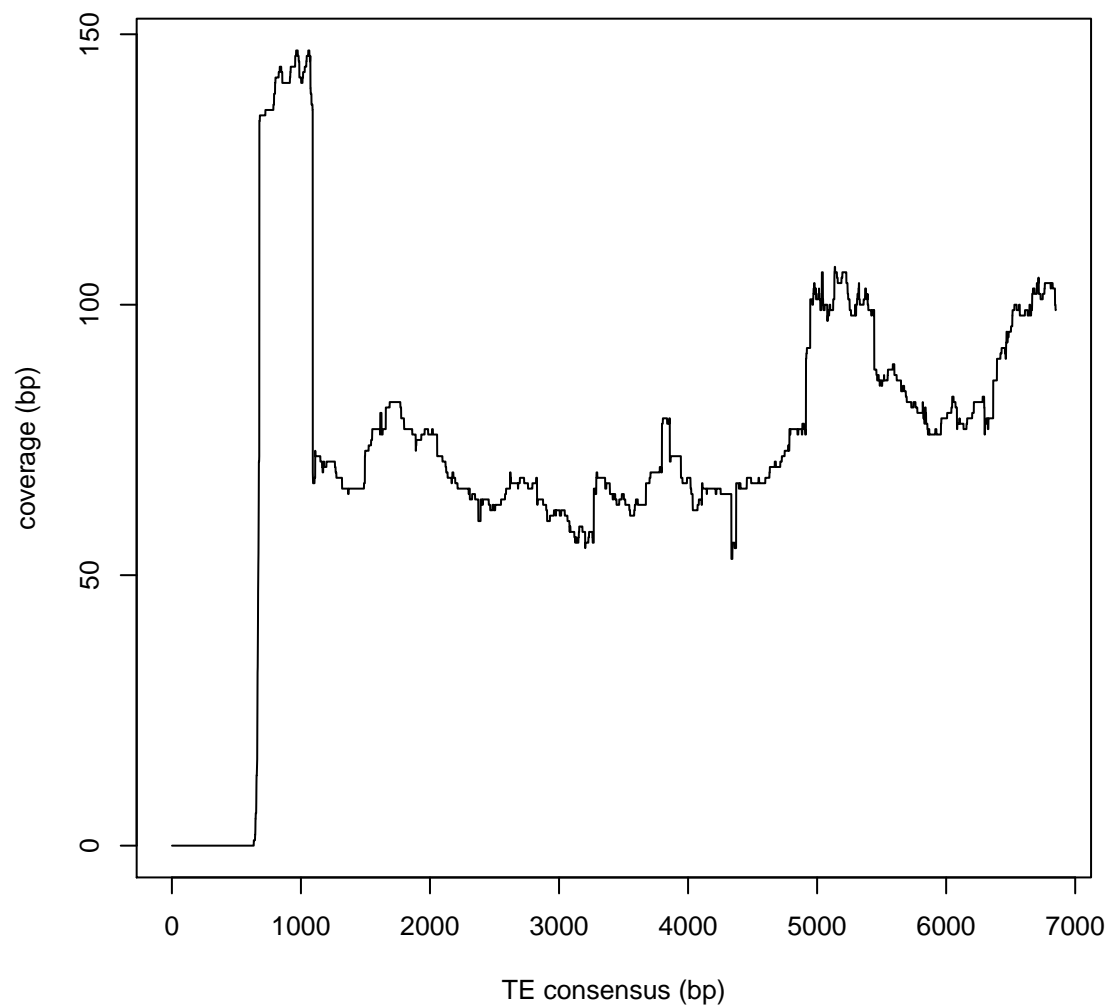


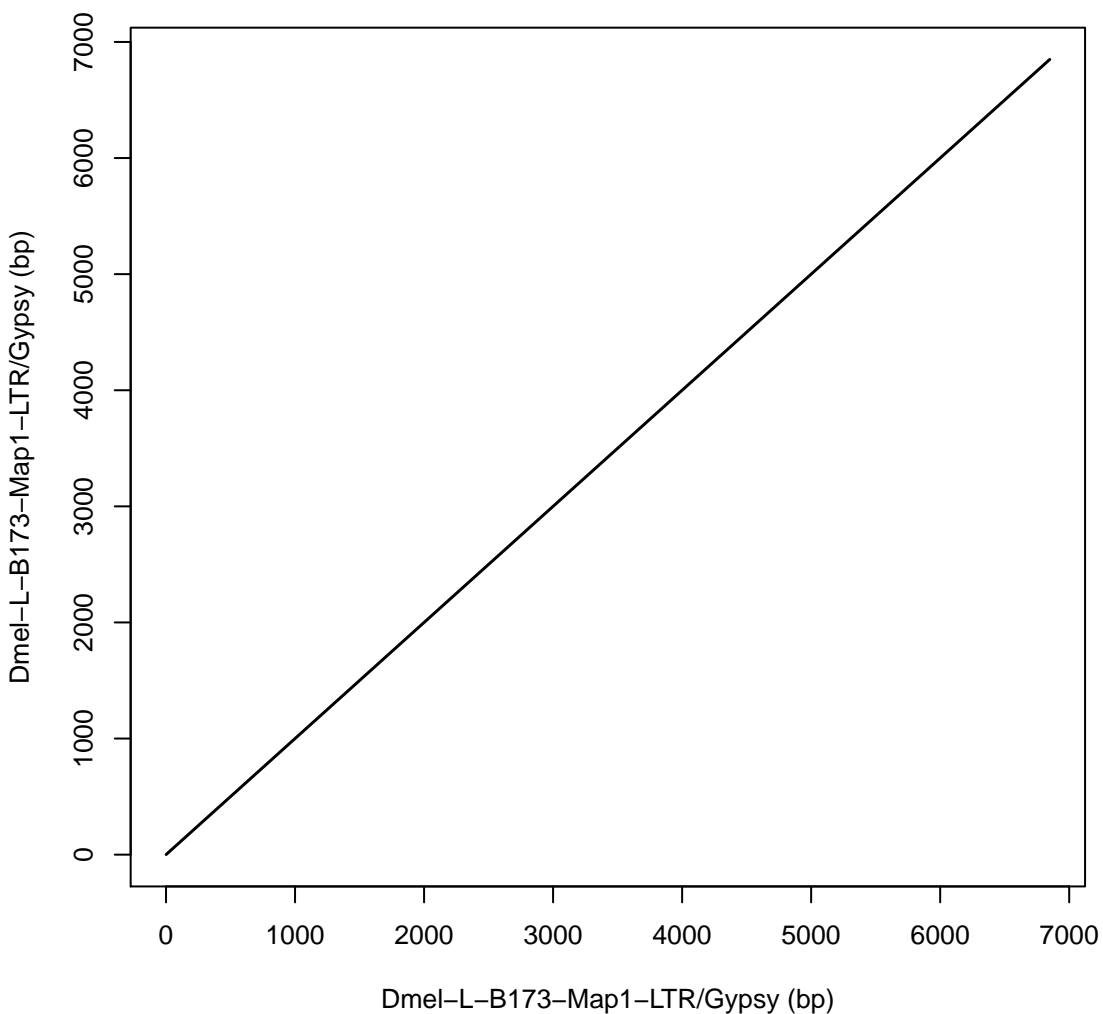
TE: Dmel-L-B173-Map1-LTR/Gypsy
consensus size: 6849bp; fragments: 459; full length: 16 (>=6164.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

