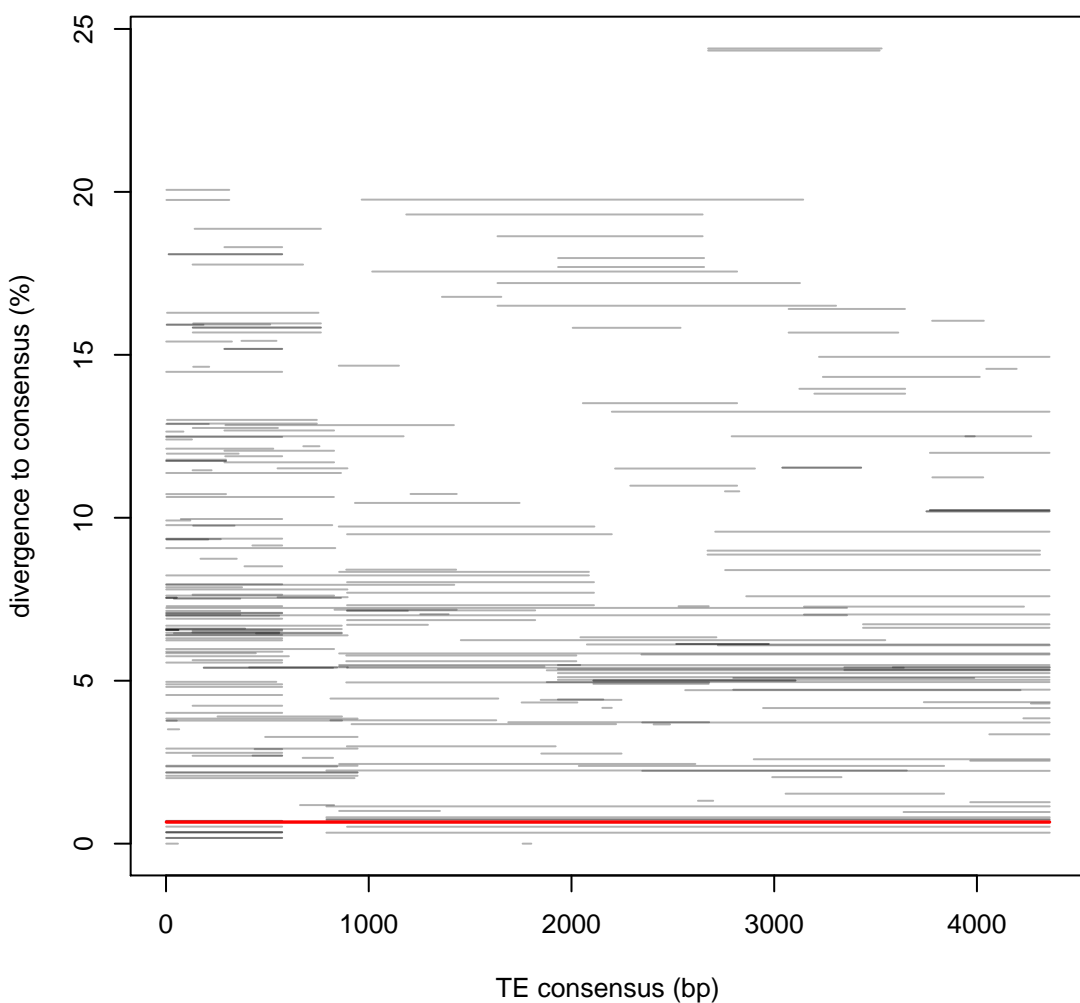
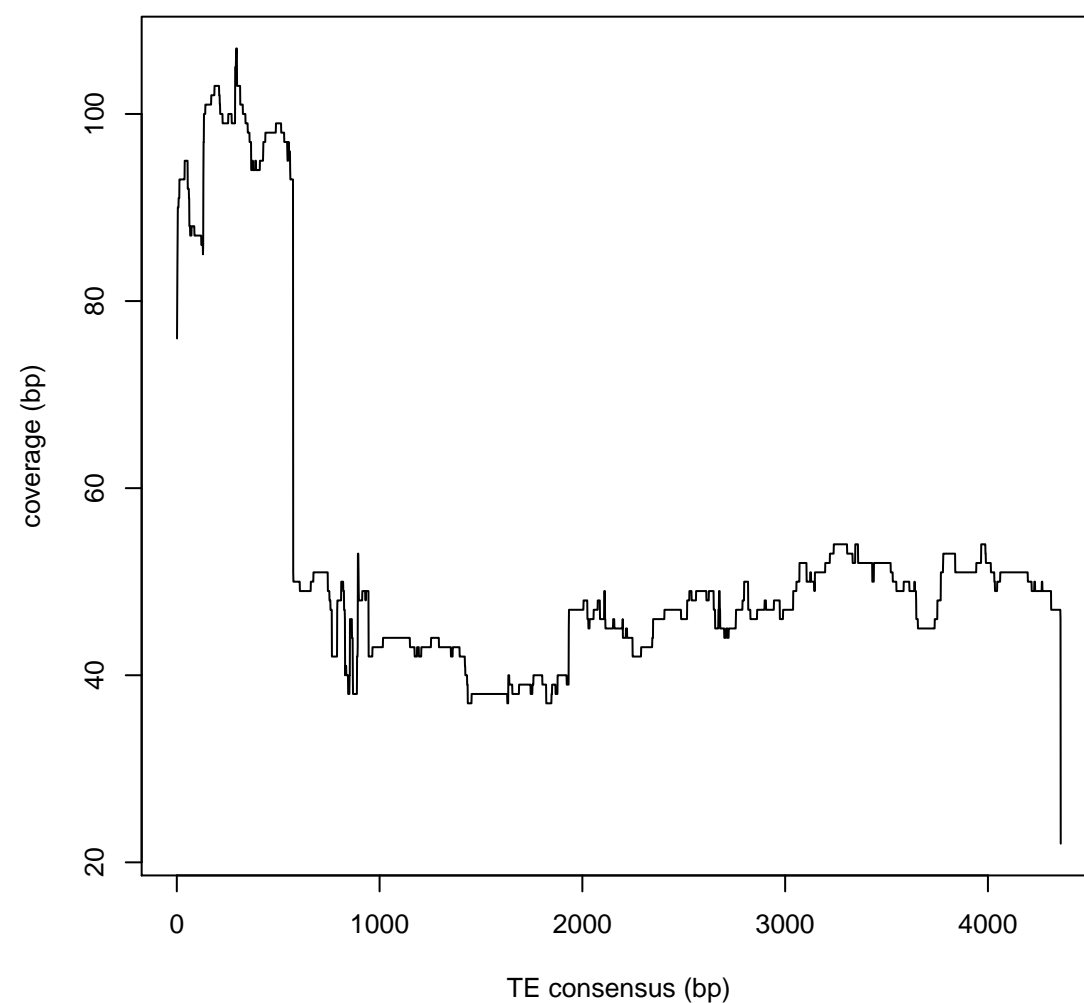


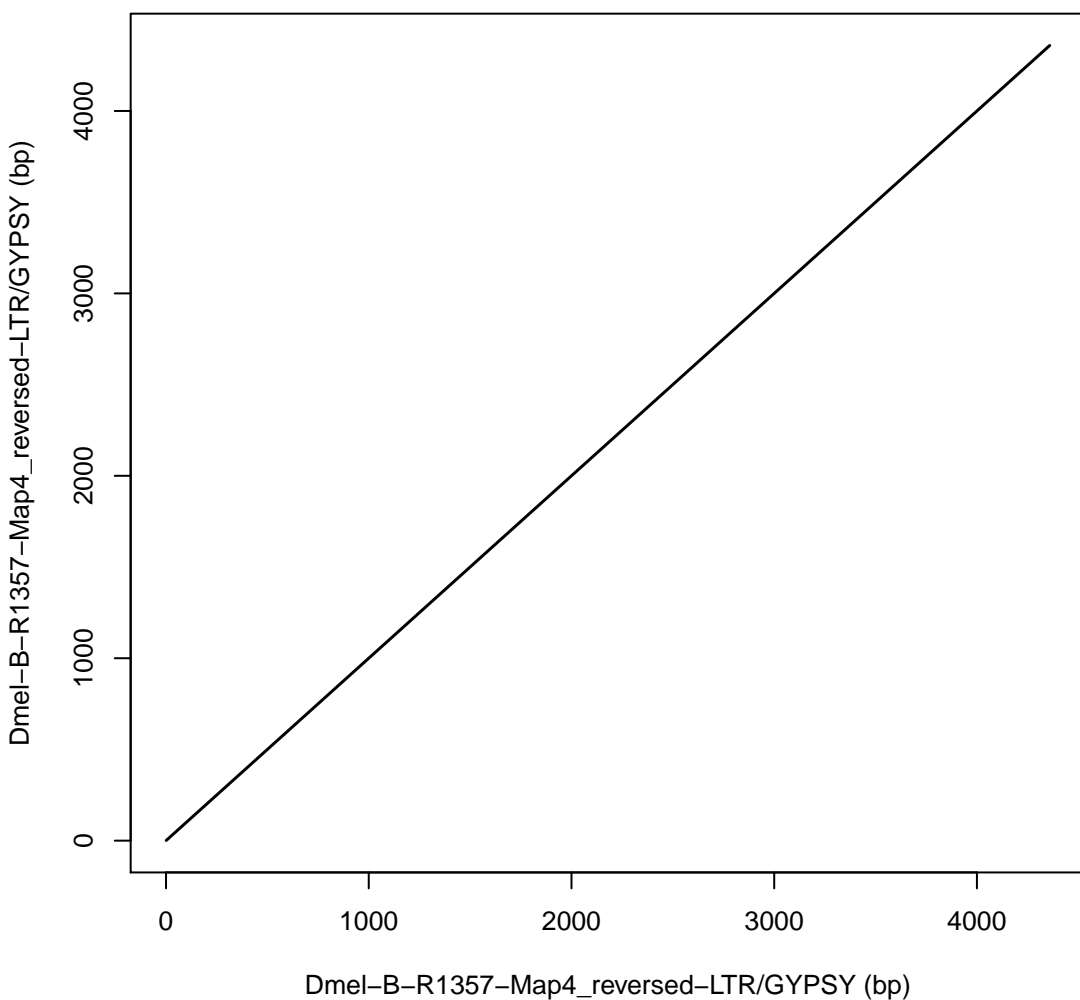
TE: Dmel-B-R1357-Map4_reversed-LTR/GYPsy
consensus size: 4359bp; fragments: 264; full length: 2 (>=3923.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

