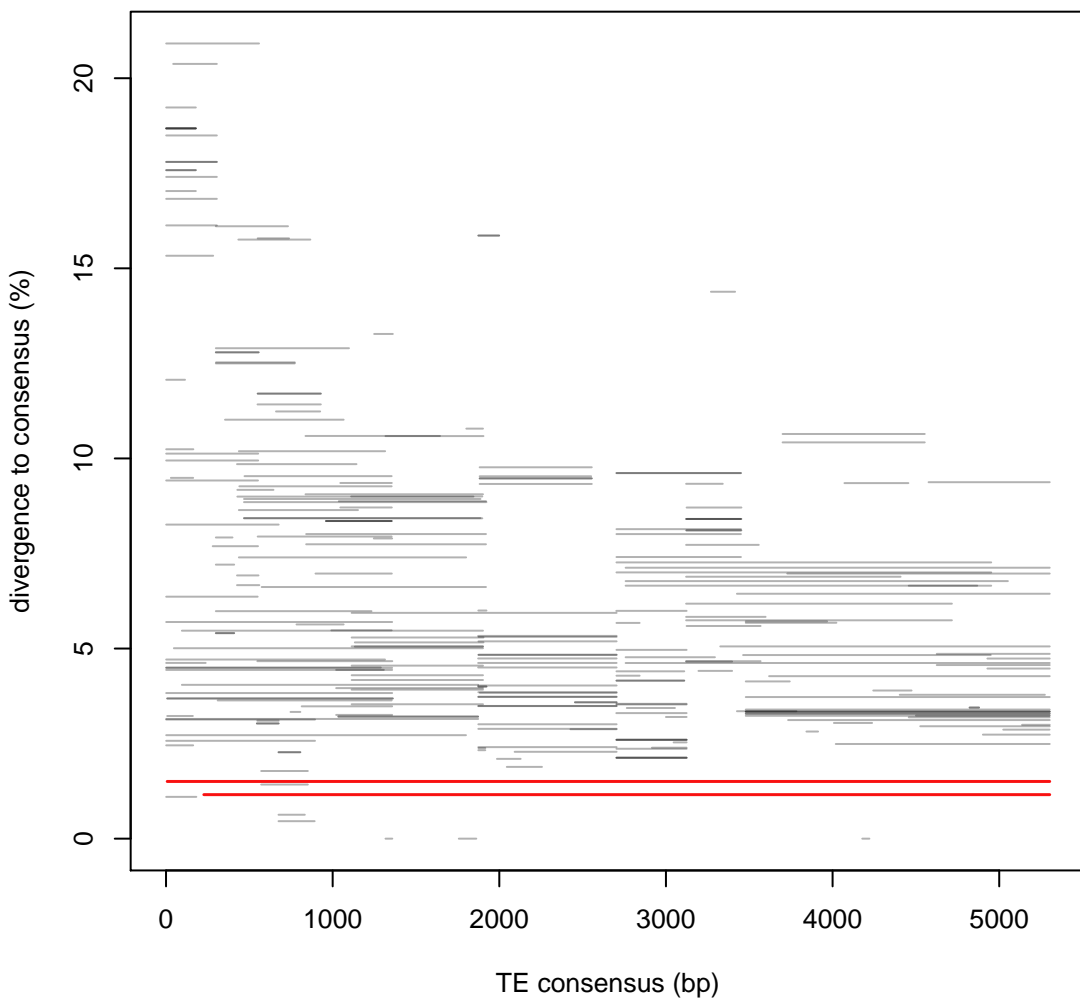
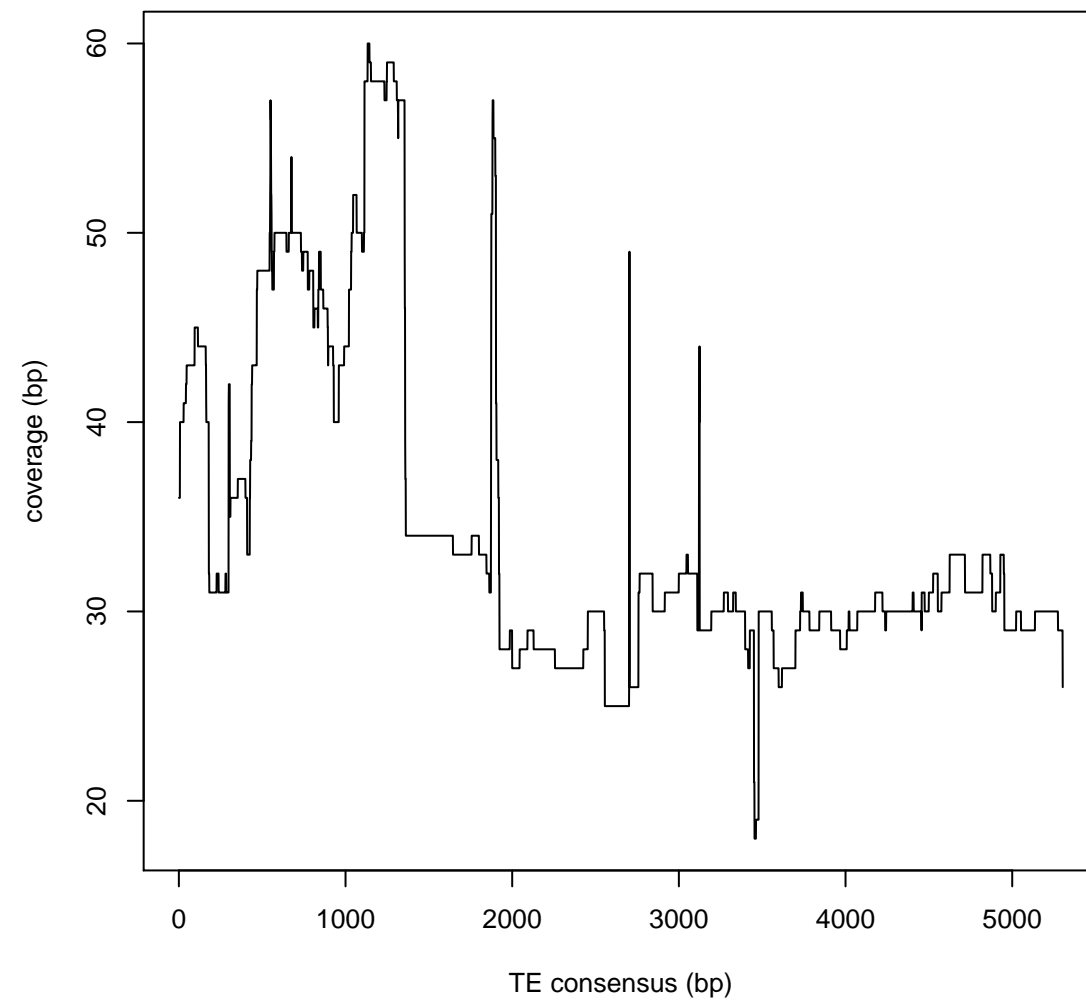


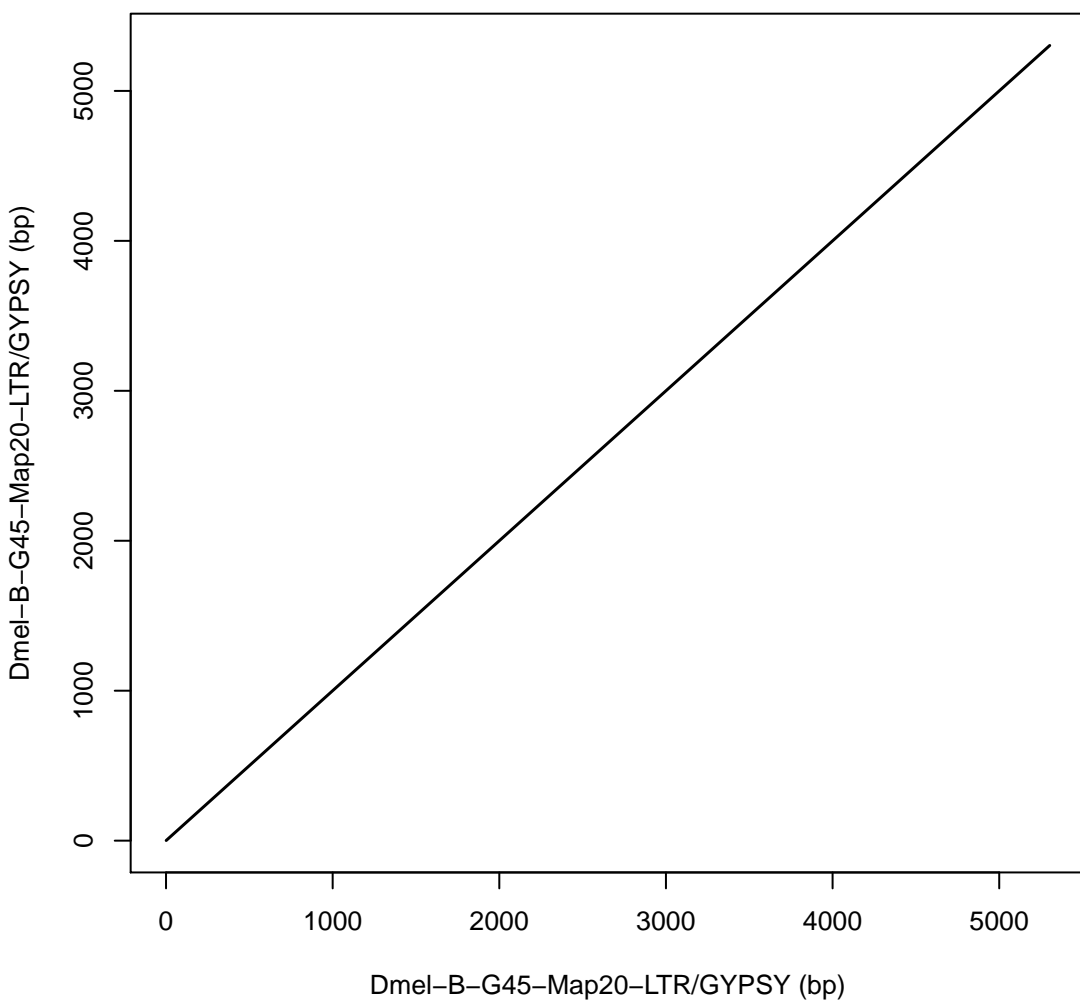
TE: Dmel-B-G45-Map20-LTR/GYPSY
consensus size: 5303bp; fragments: 256; full length: 2 (>=4772.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

