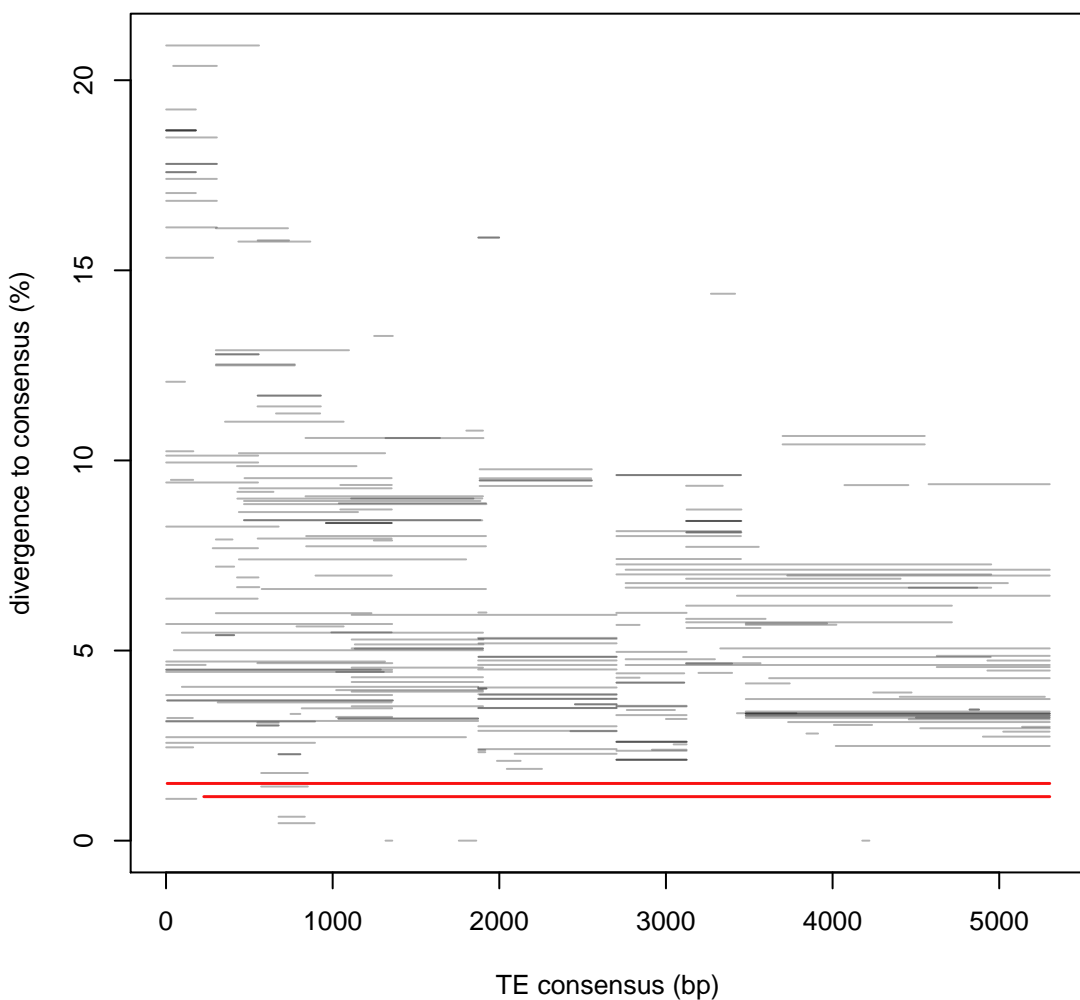
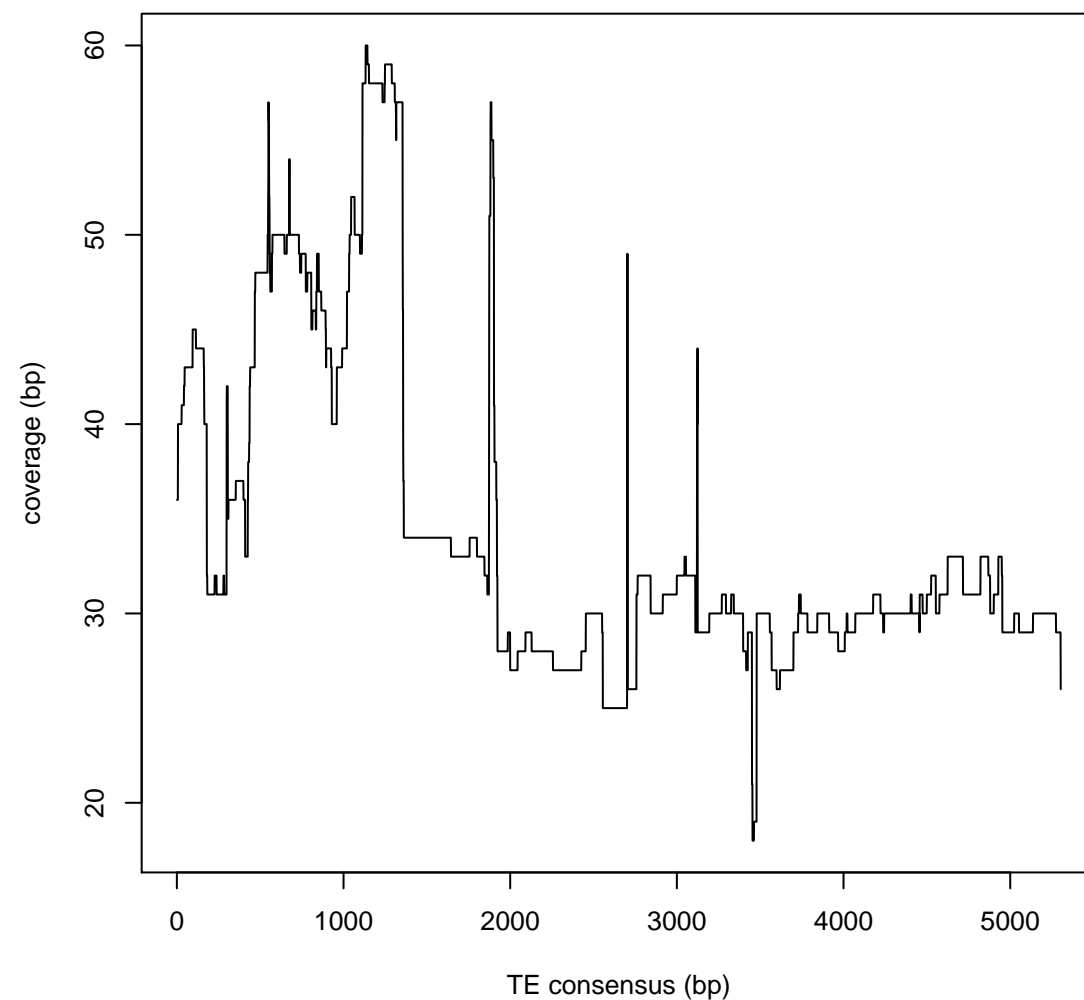


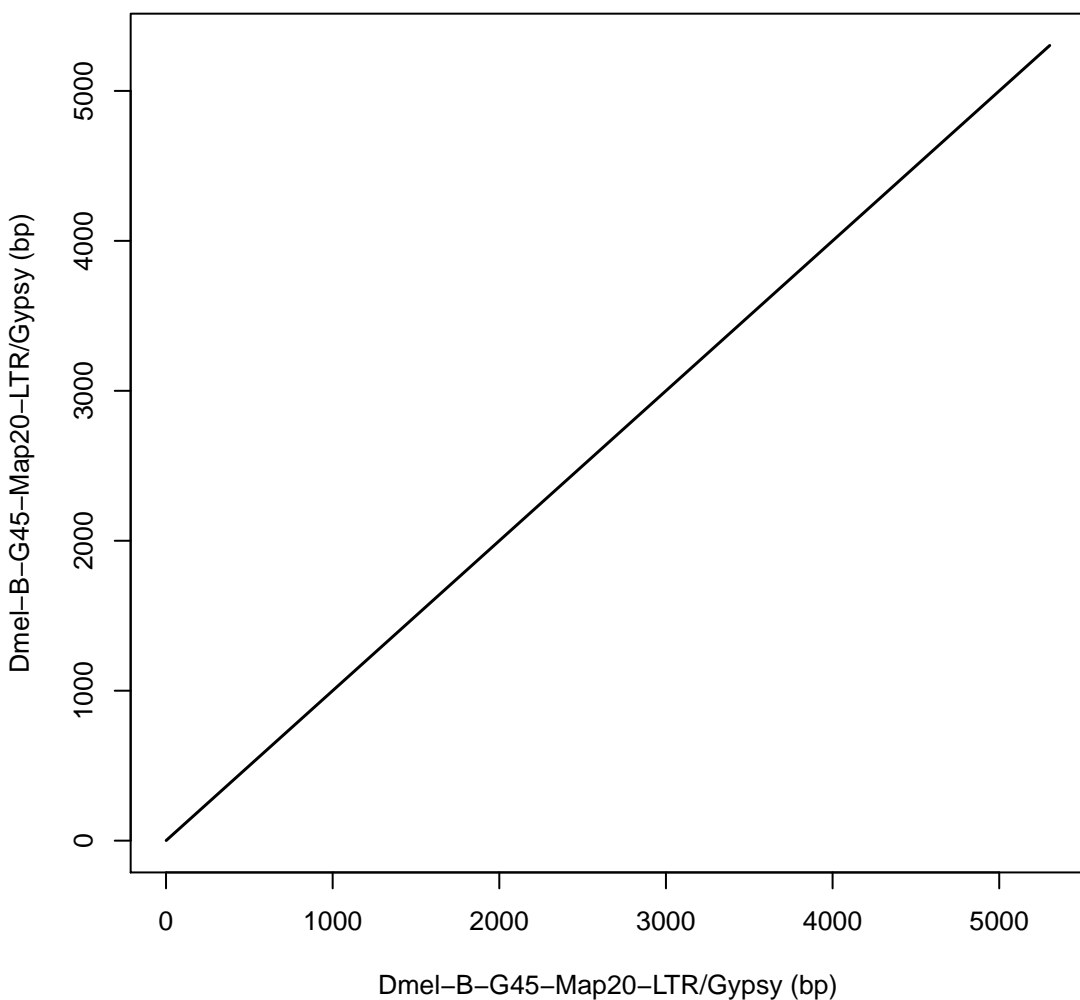
TE: Dmel-B-G45-Map20-LTR/Gypsy
consensus size: 5303bp; fragments: 256; full length: 2 (≥ 4772.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

