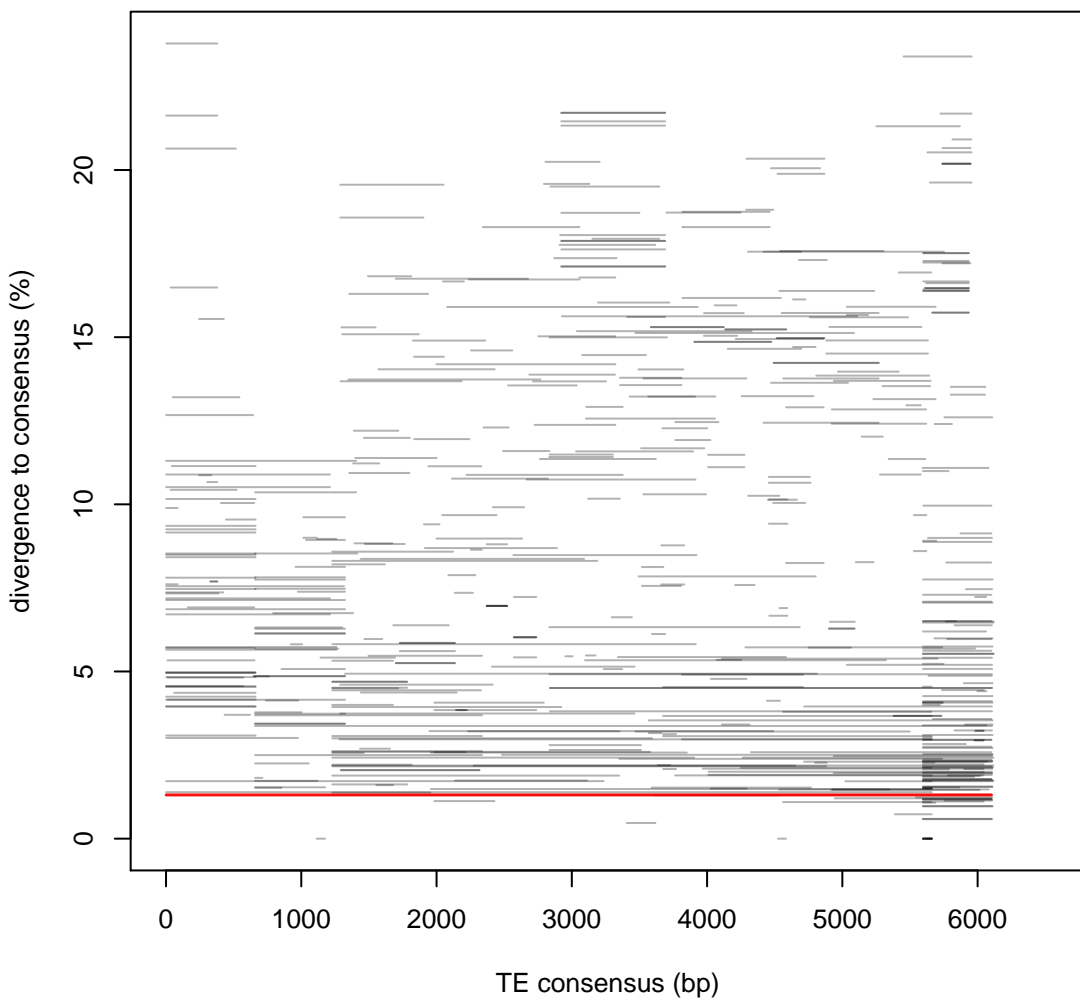
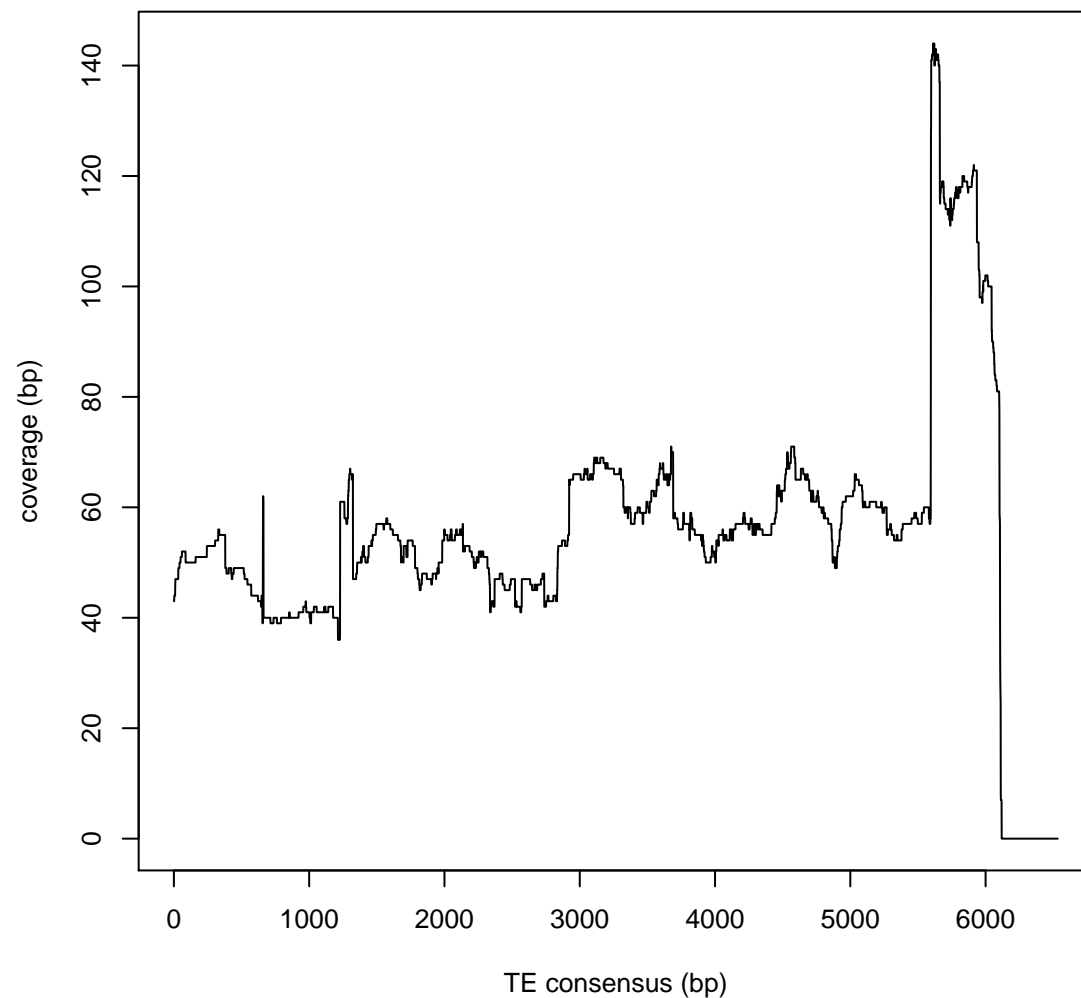


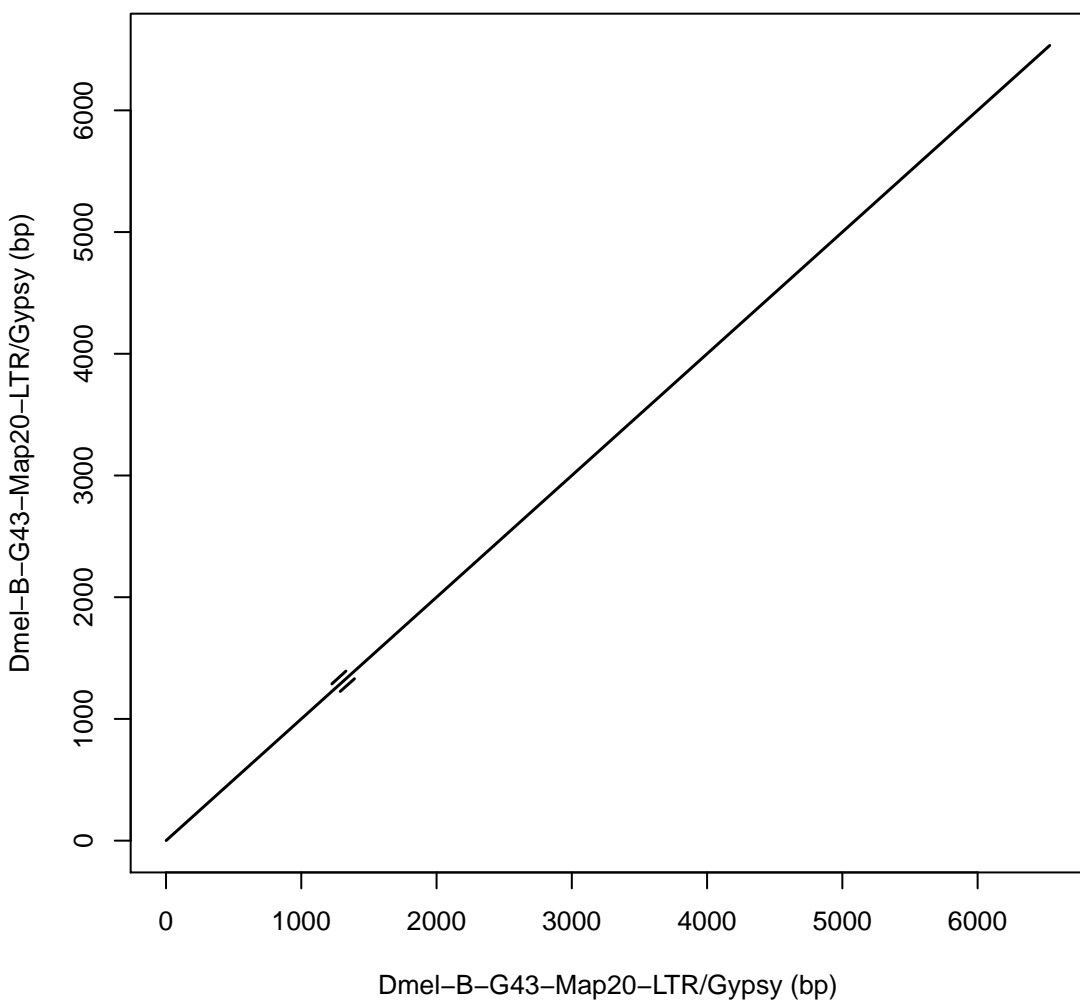
TE: Dmel-B-G43-Map20-LTR/Gypsy
consensus size: 6533bp; fragments: 523; full length: 1 (≥ 5879.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

