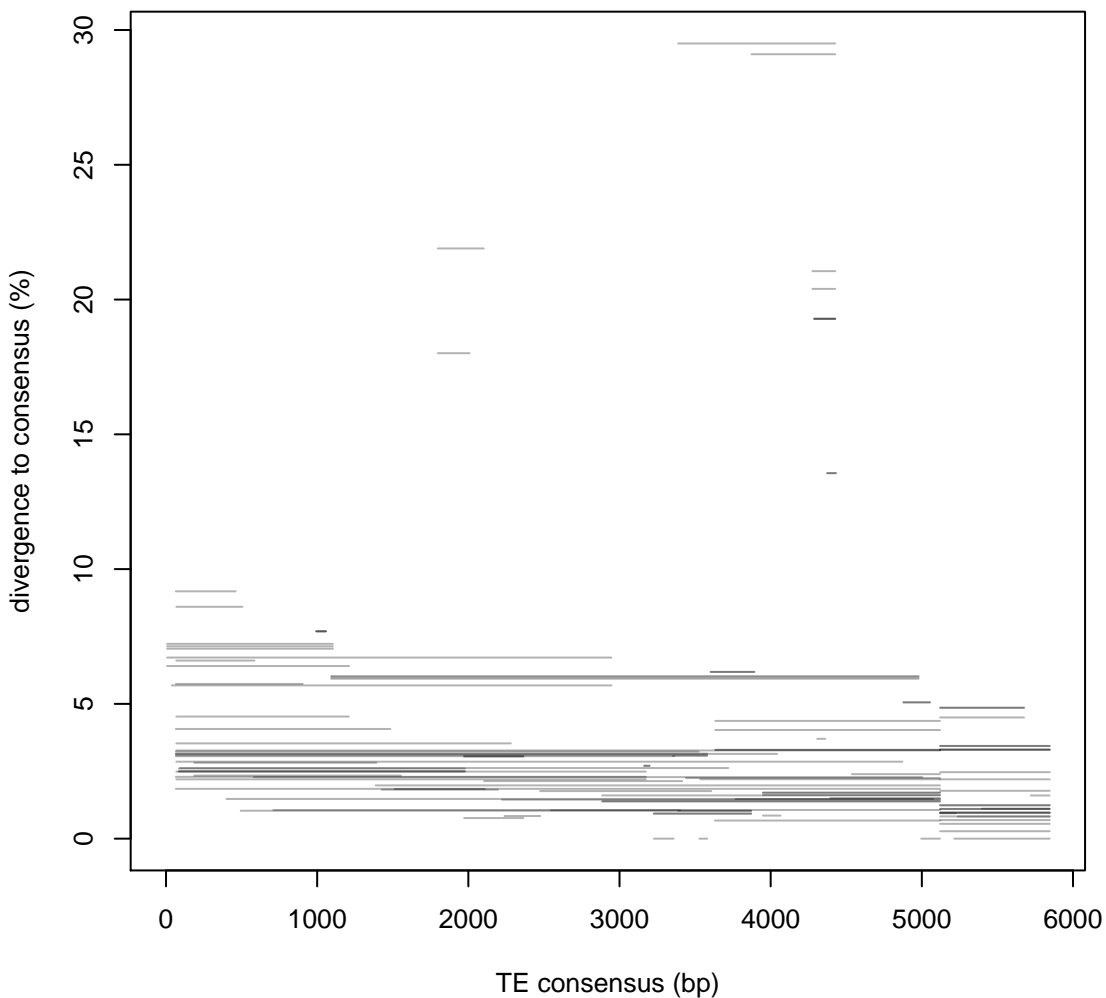
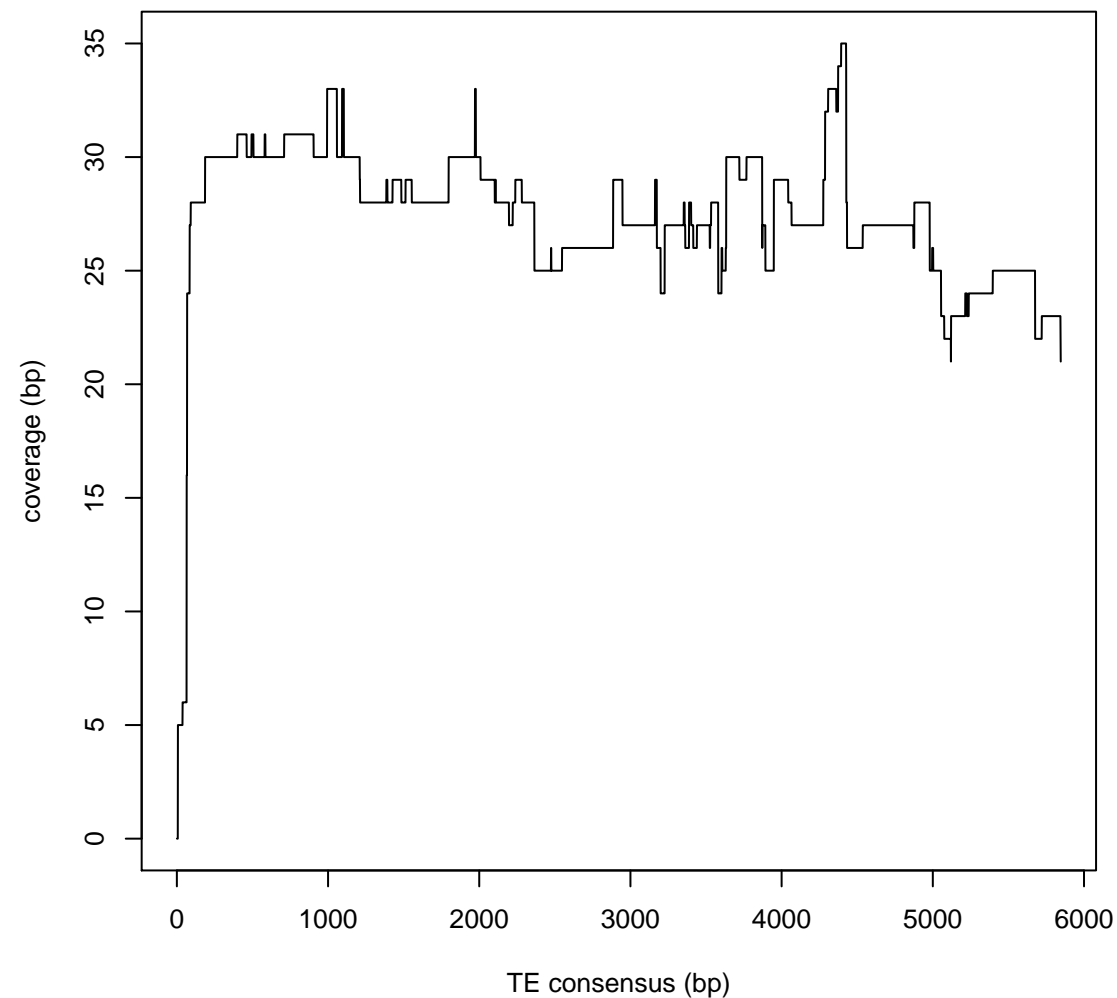


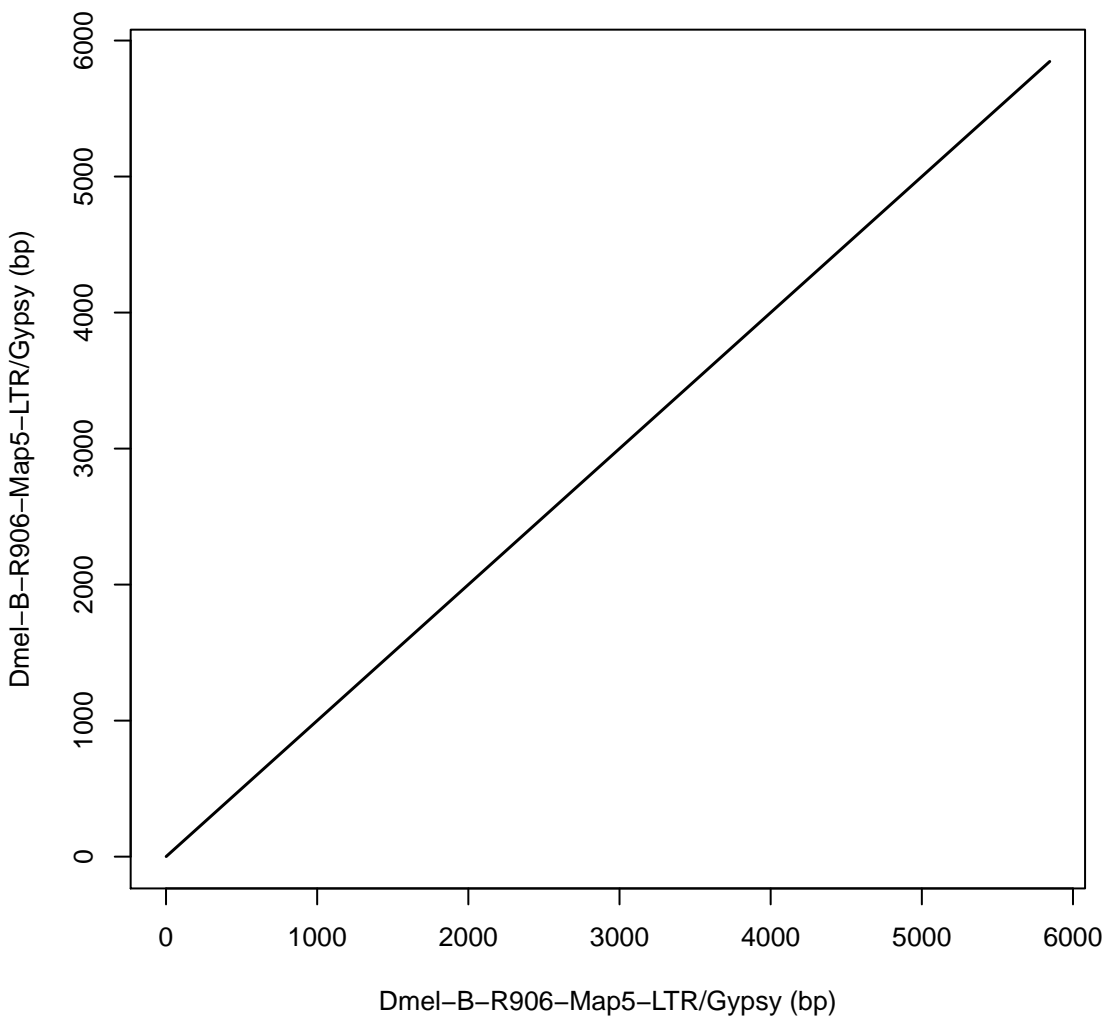
TE: Dmel-B-R906-Map5-LTR/Gypsy
consensus size: 5846bp; fragments: 120; full length: 0 (>=5261.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

