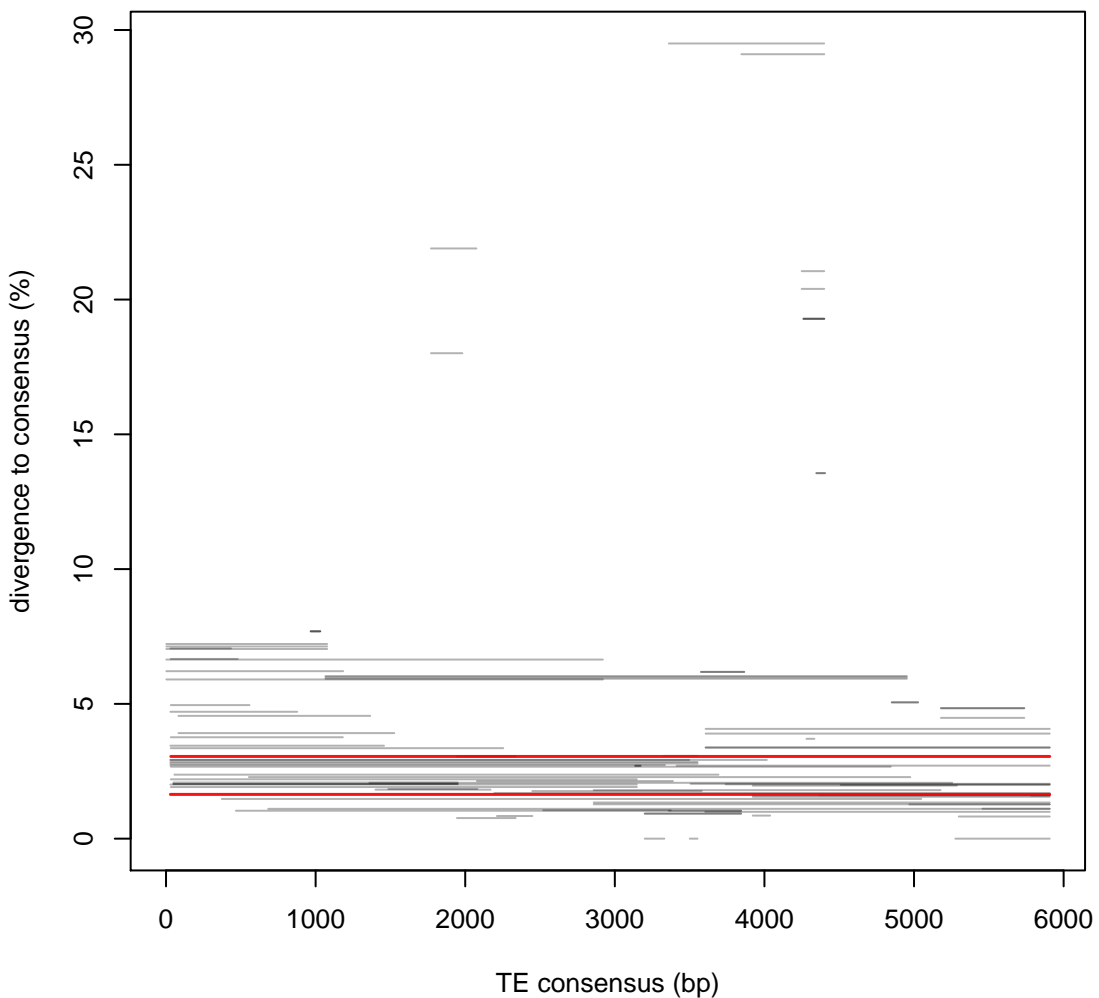
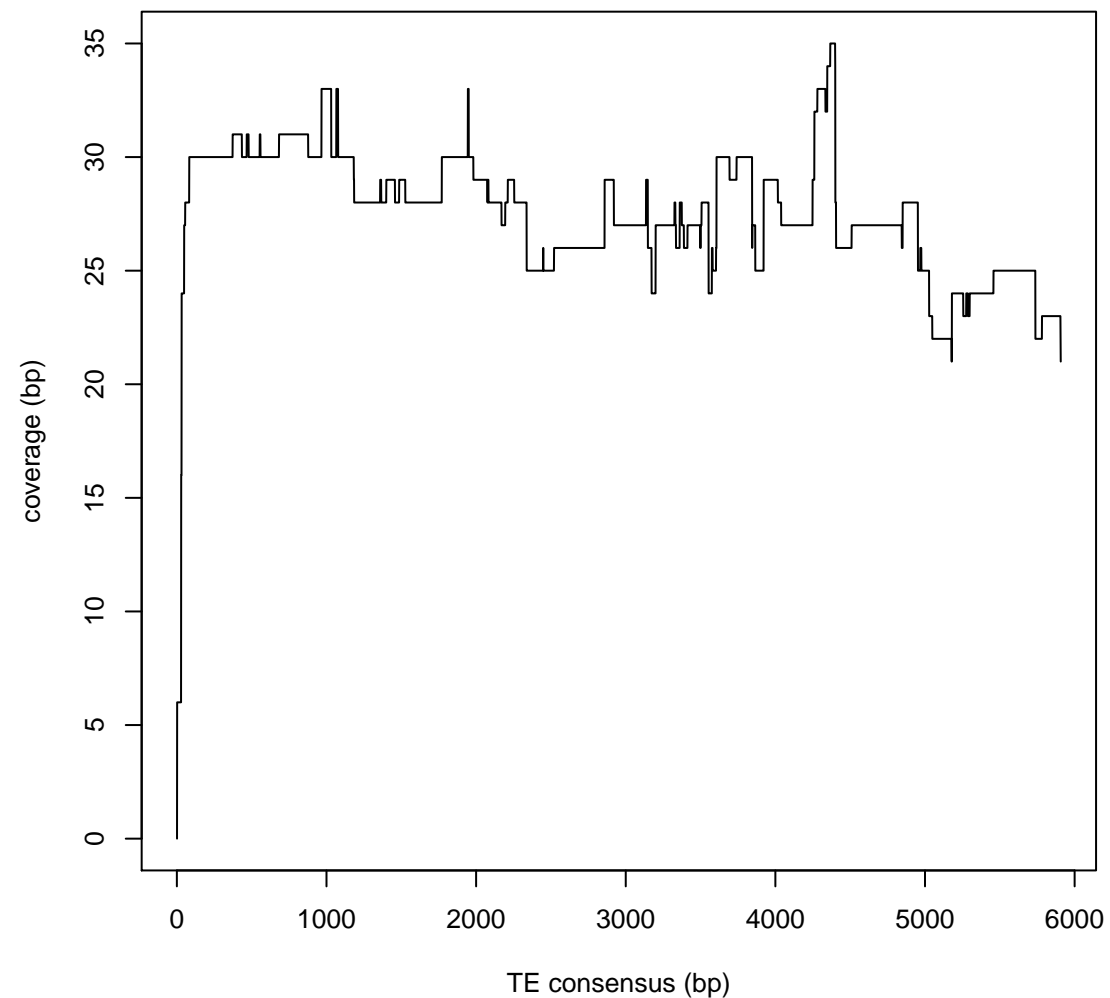


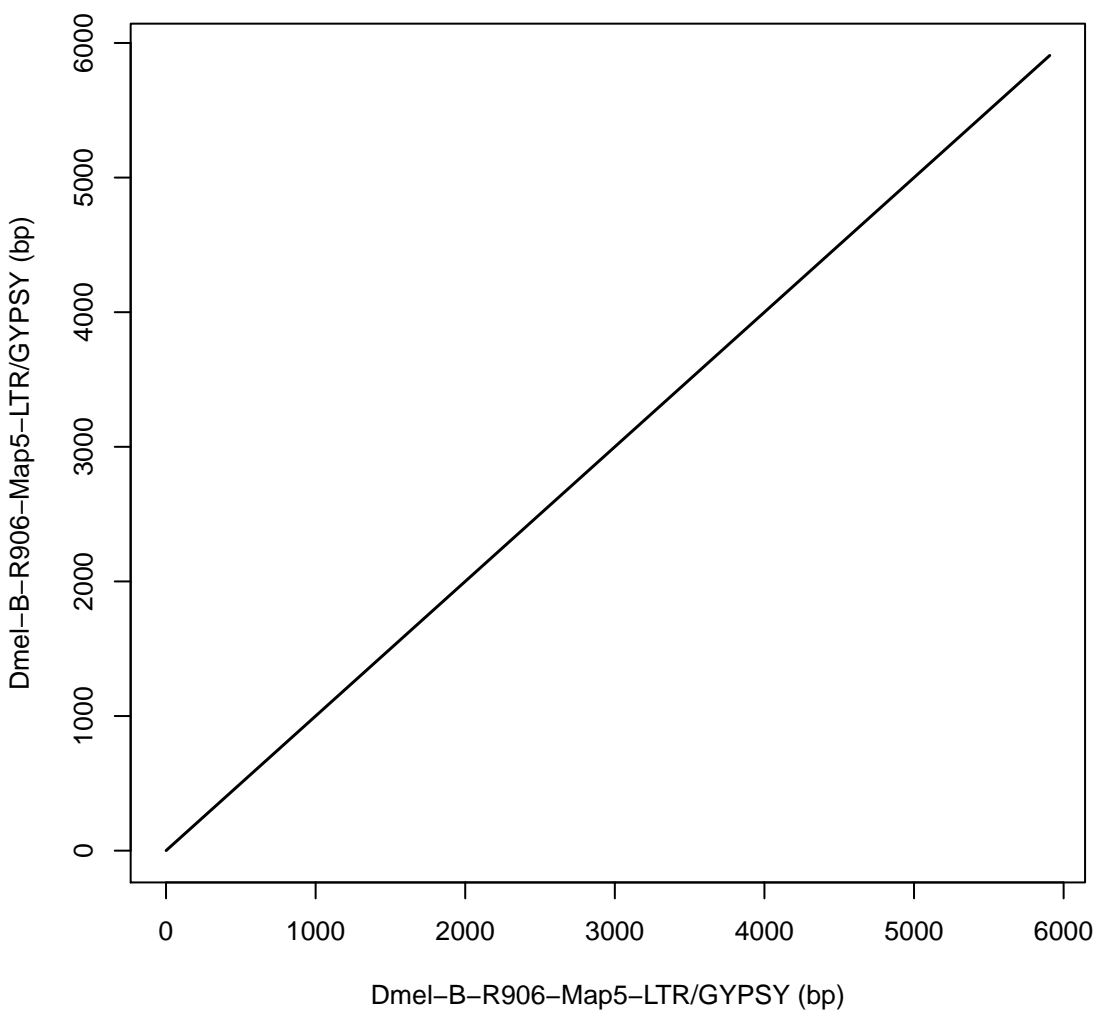
TE: Dmel-B-R906-Map5-LTR/GYPSY
consensus size: 5907bp; fragments: 100; full length: 2 (>=5316.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

