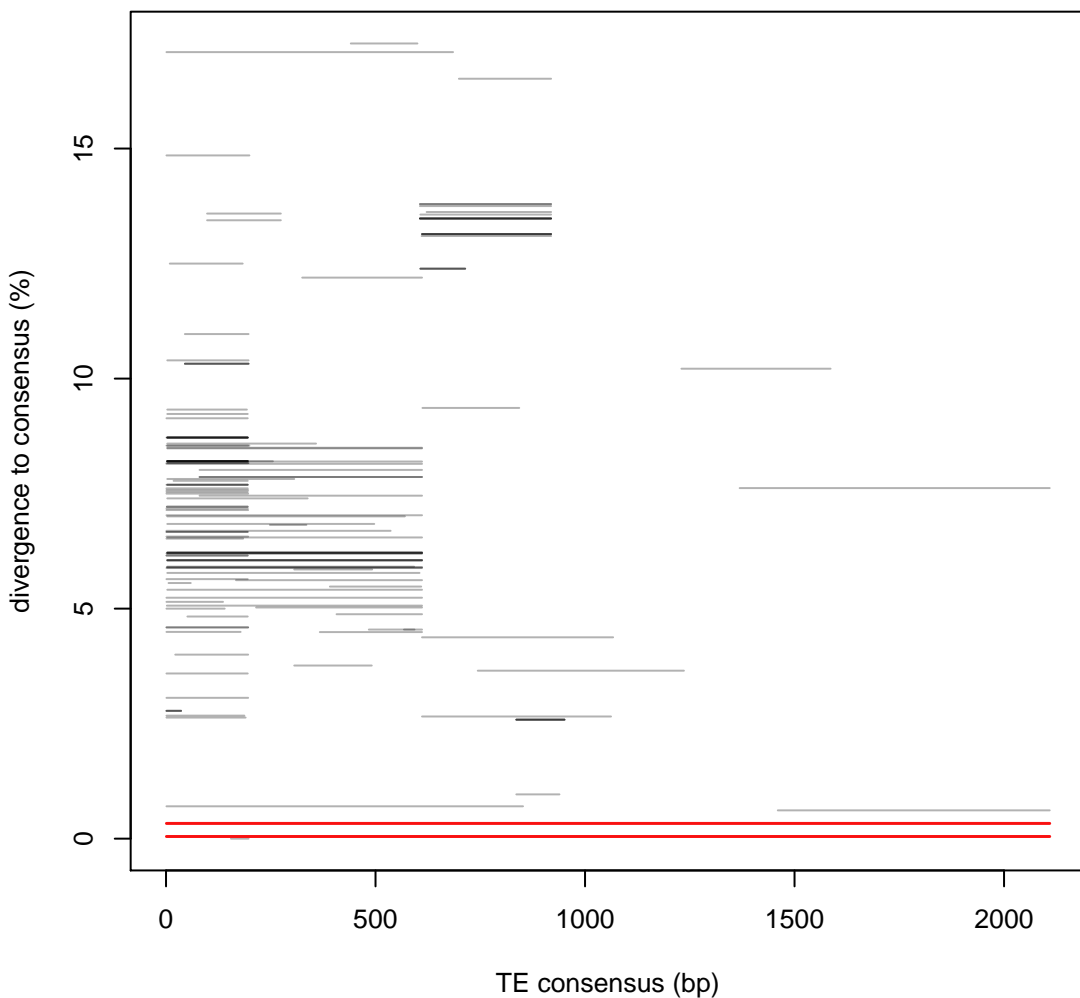
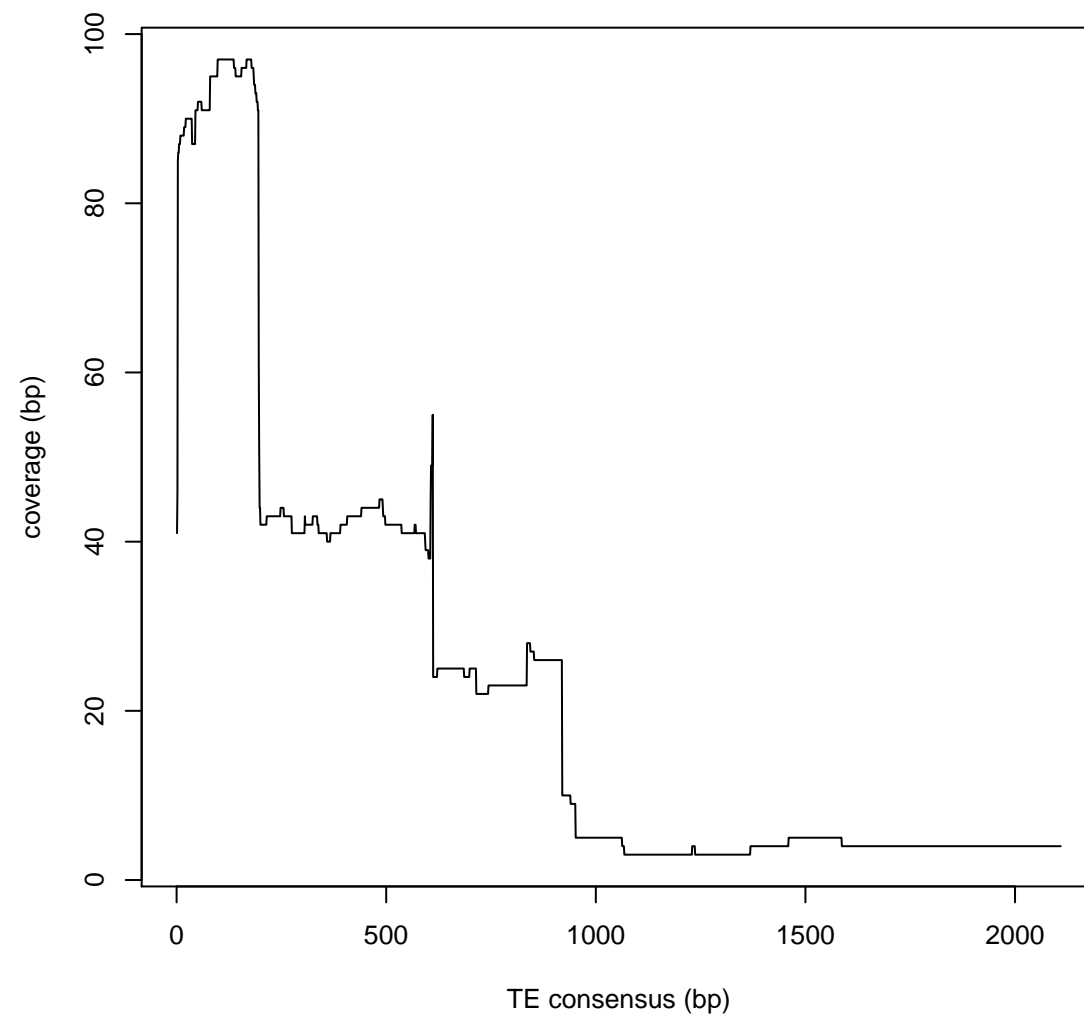


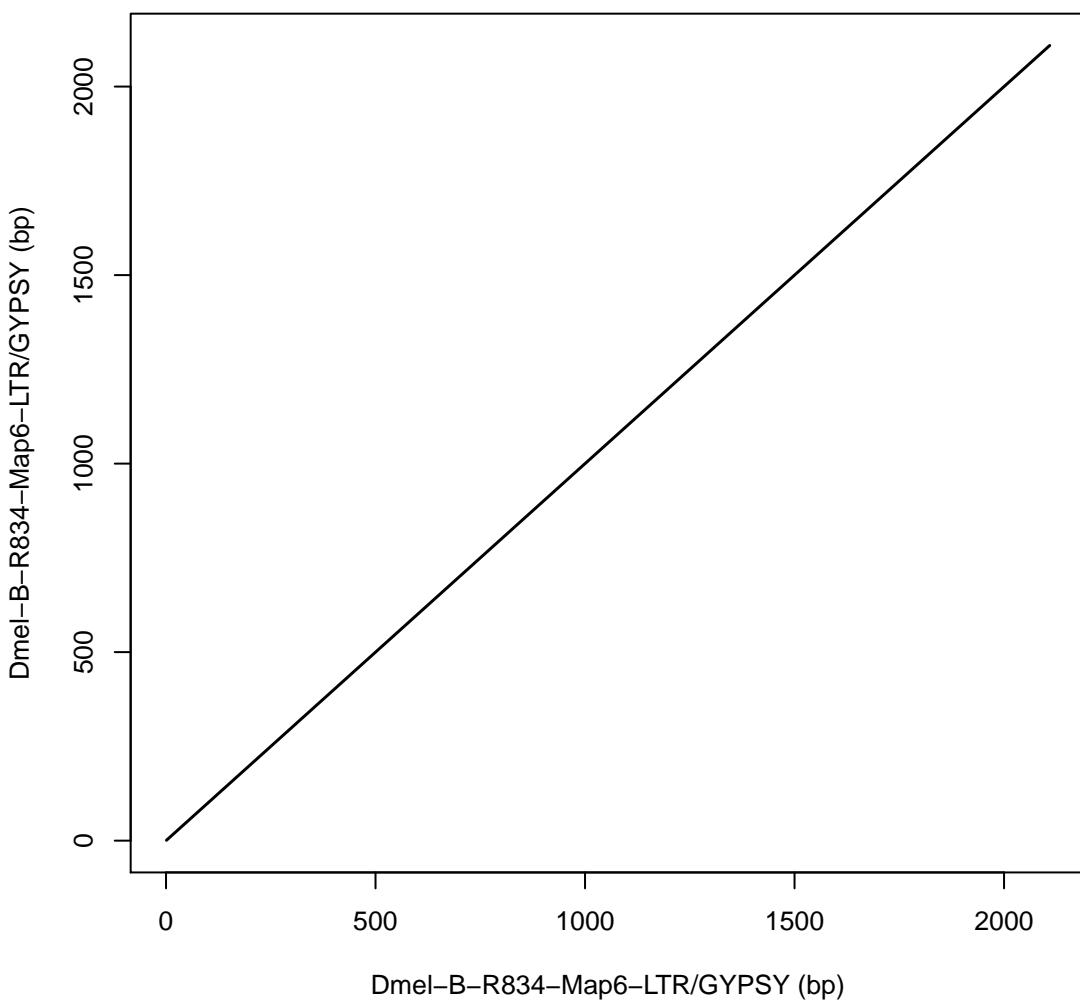
TE: Dmel-B-R834-Map6-LTR/GYPSY
consensus size: 2109bp; fragments: 145; full length: 2 (>=1898.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

