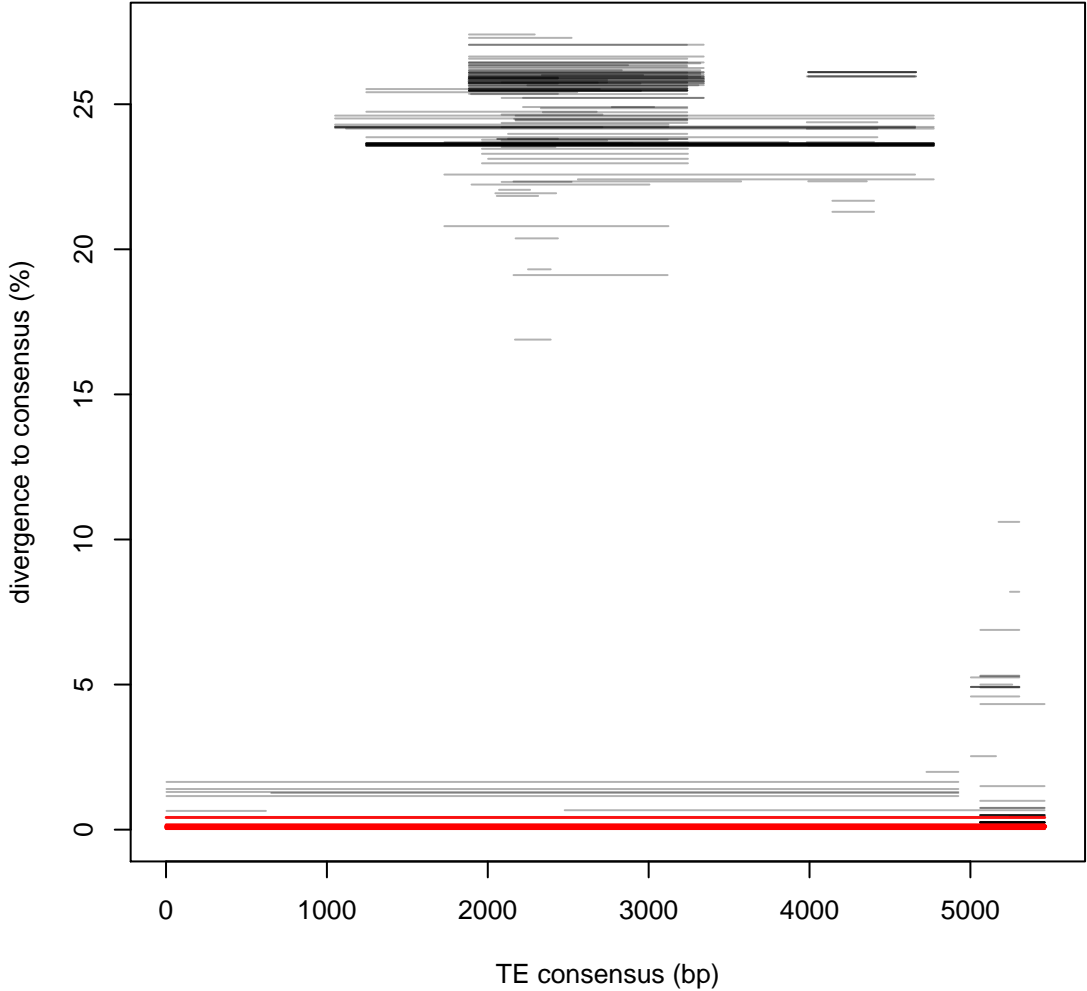
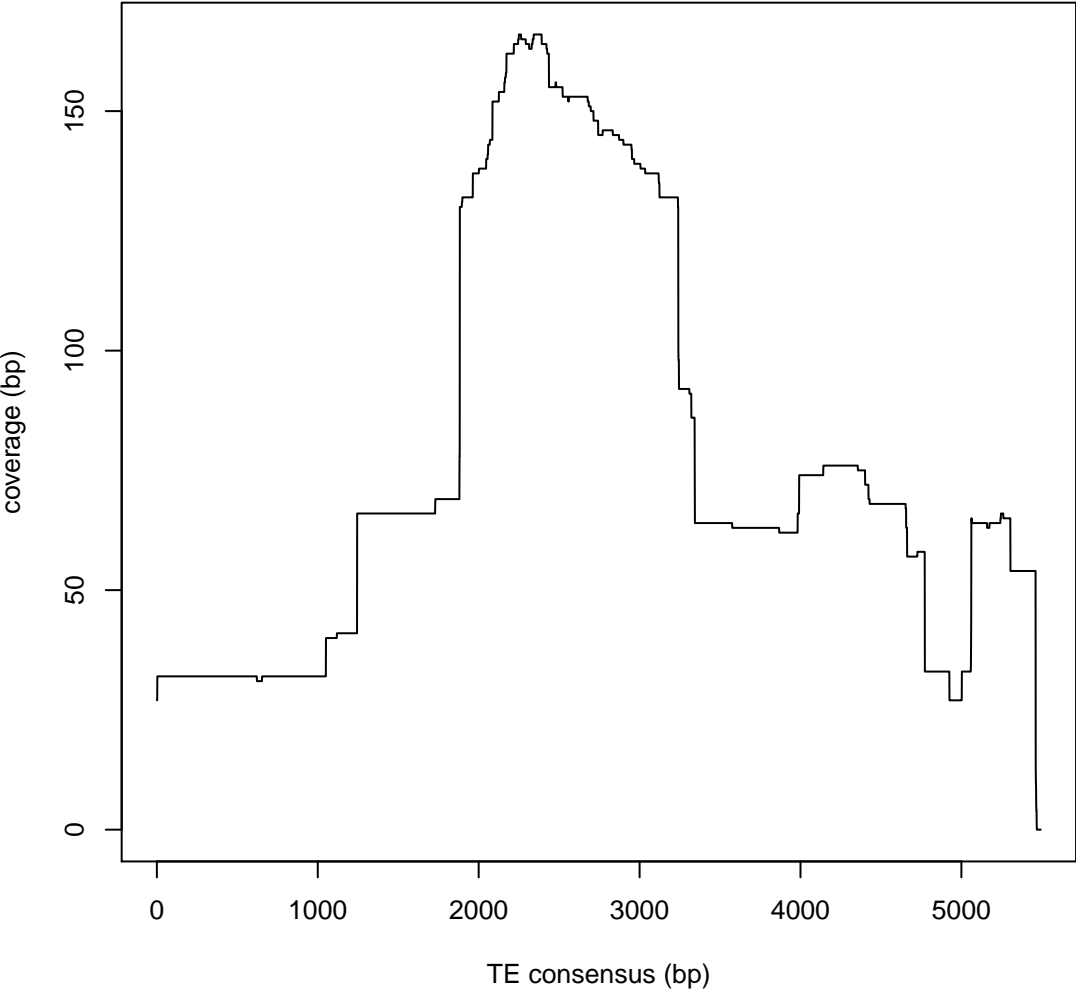


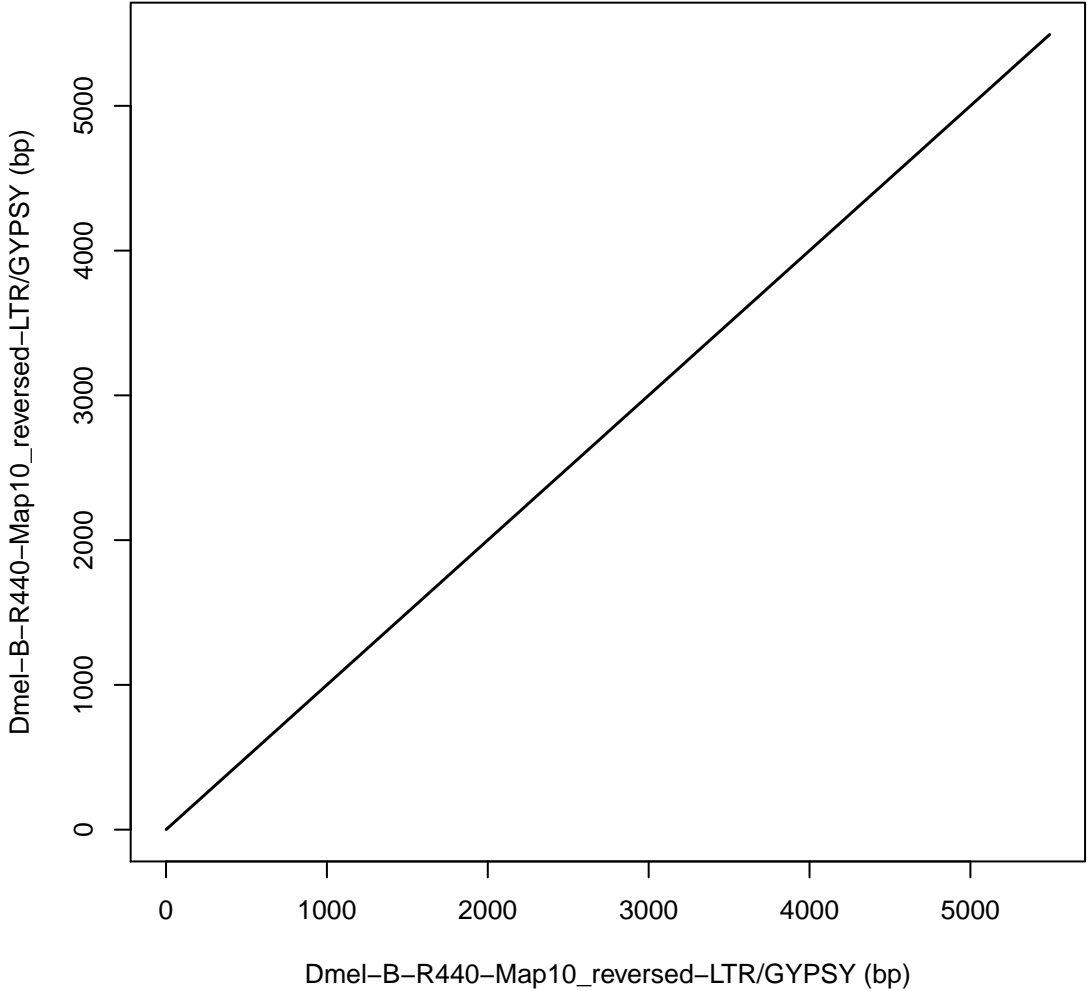
TE: Dmel-B-R440-Map10_reversed-LTR/GYPSY
consensus size: 5493bp; fragments: 229; full length: 26 (>=4943.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

