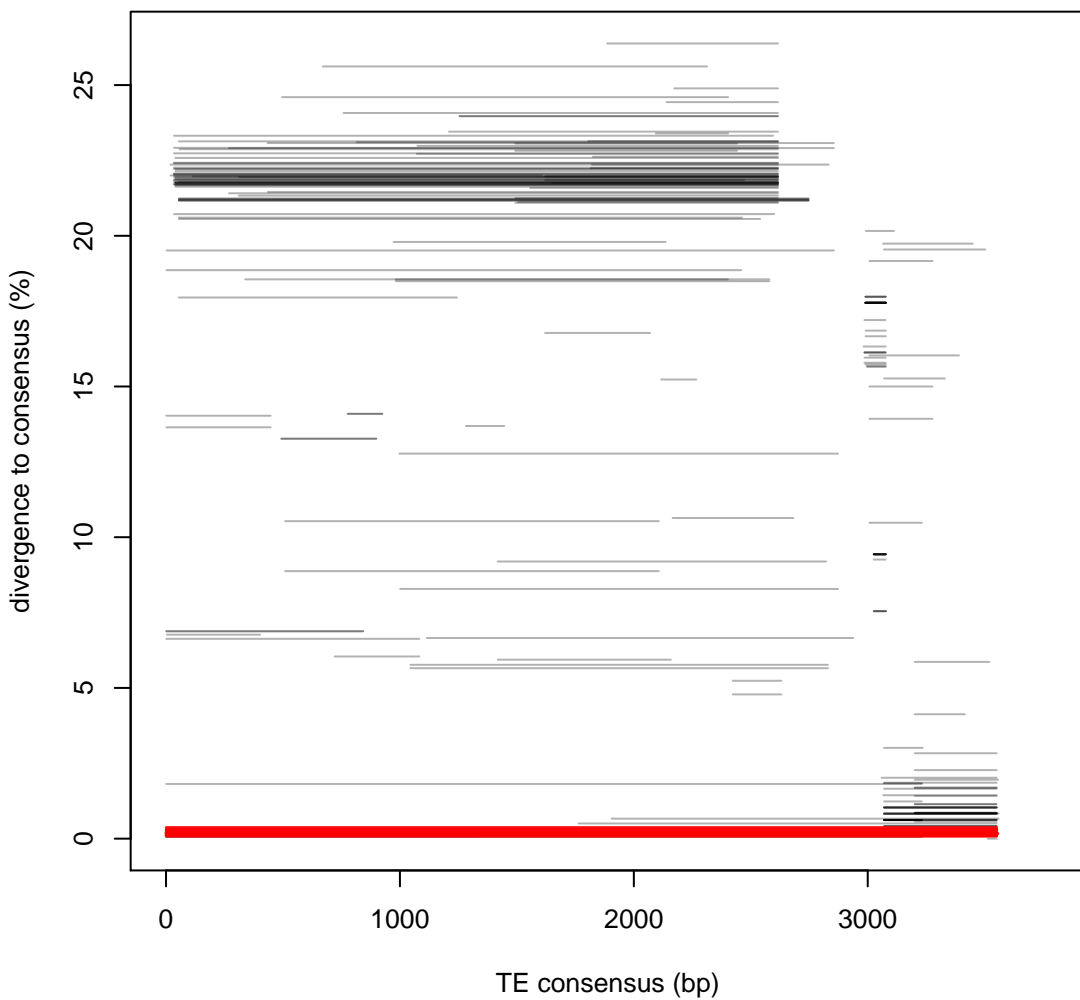
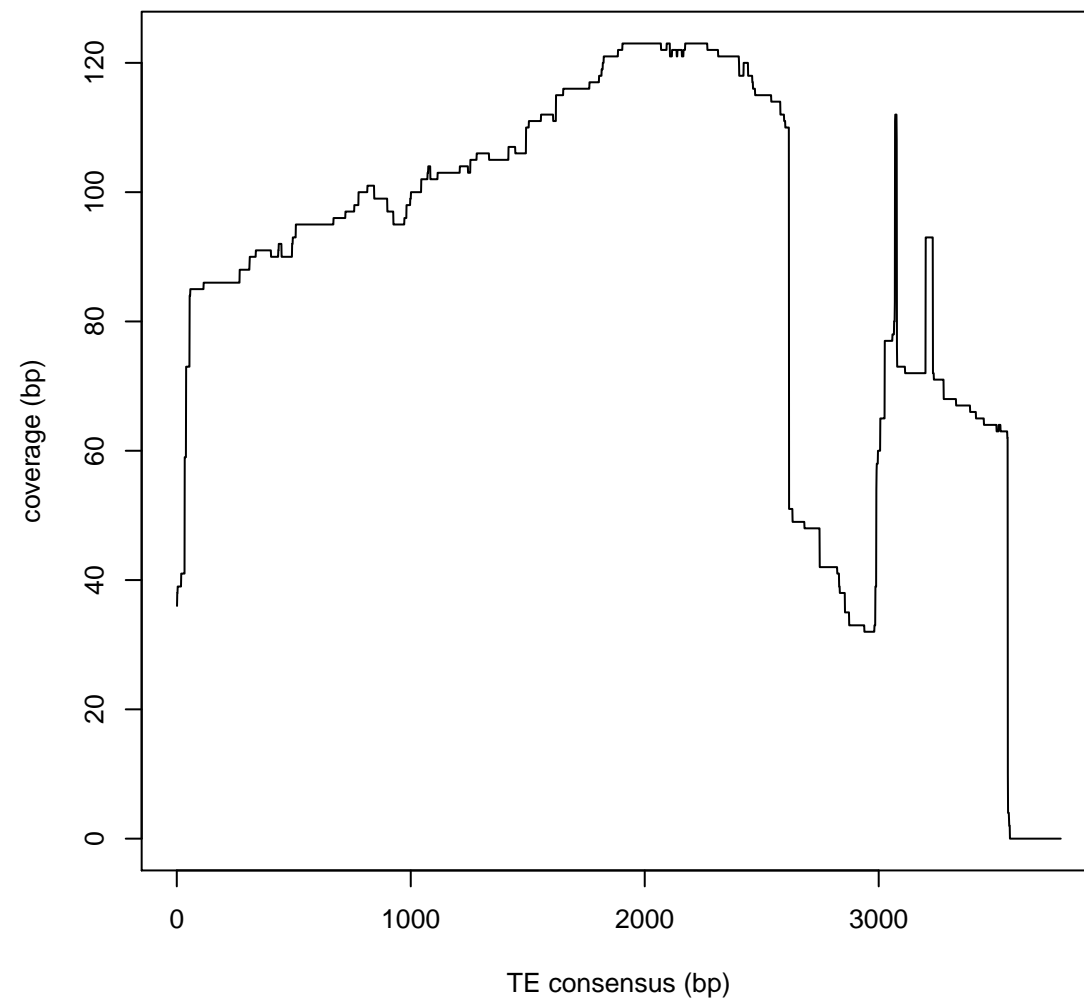


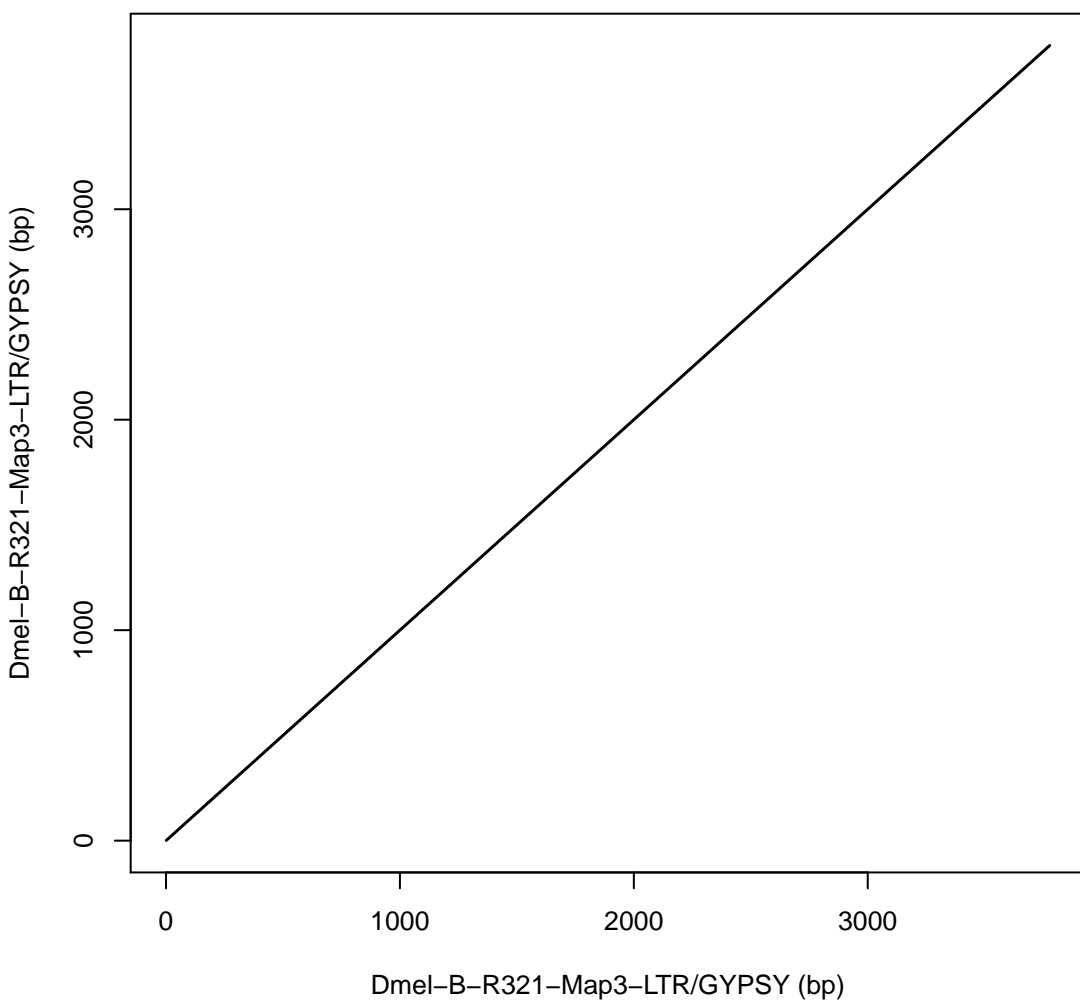
TE: Dmel-B-R321-Map3-LTR/GYPsy
consensus size: 3778bp; fragments: 247; full length: 19 (>=3400.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

