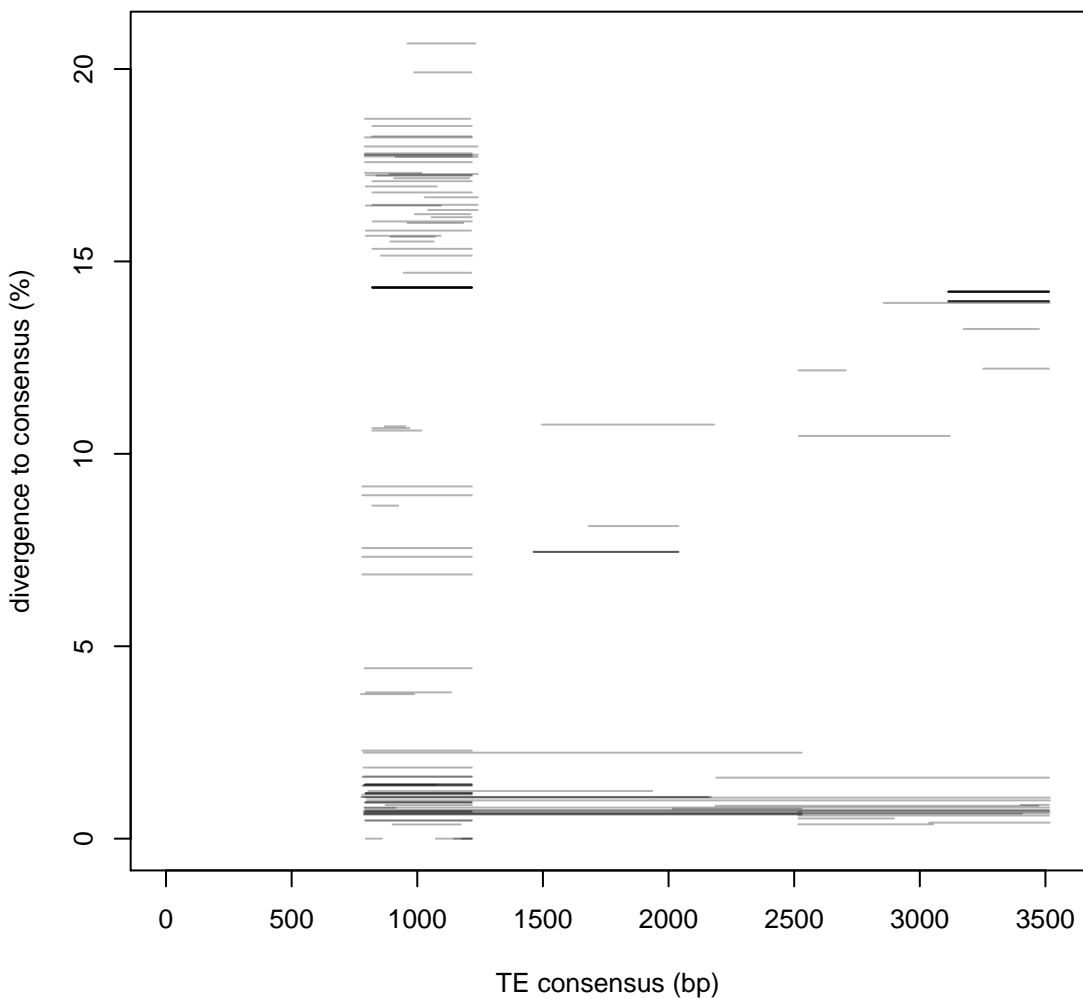
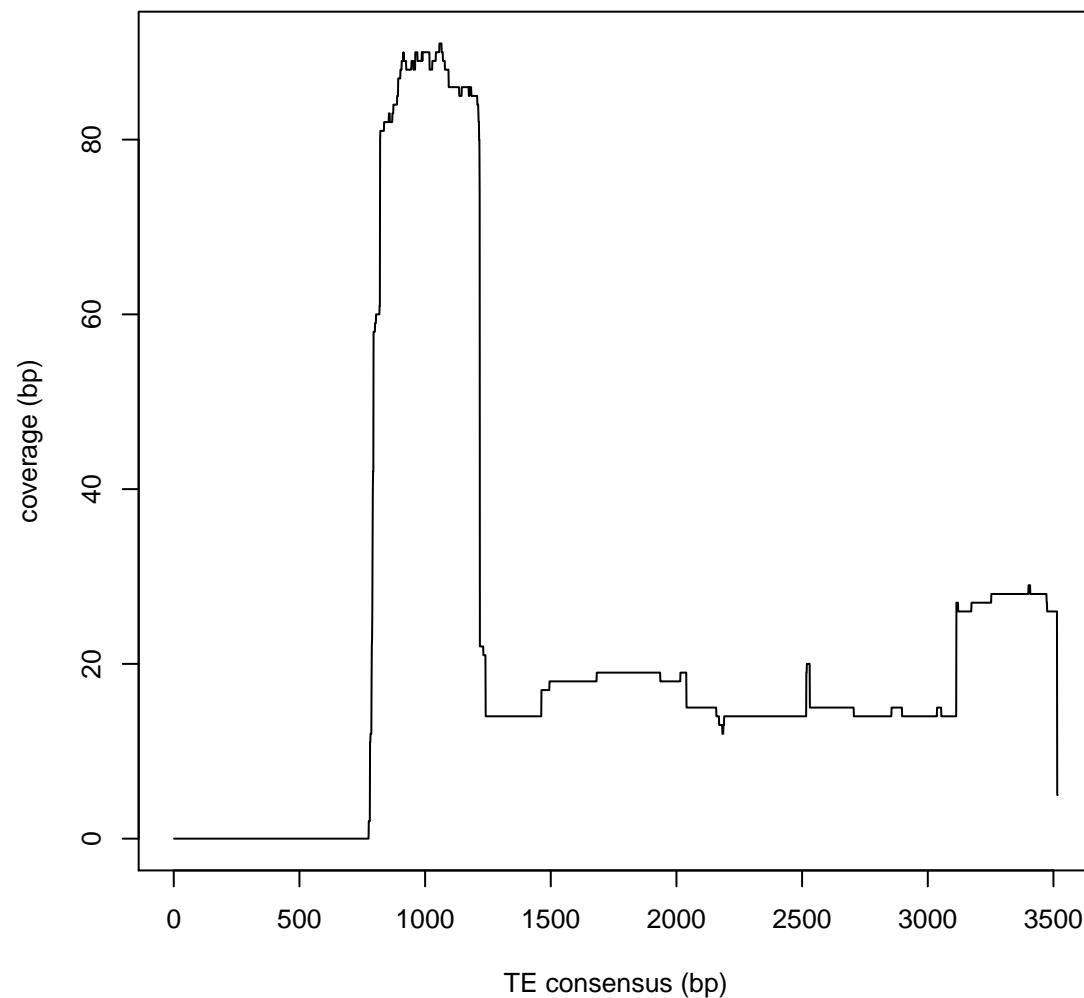


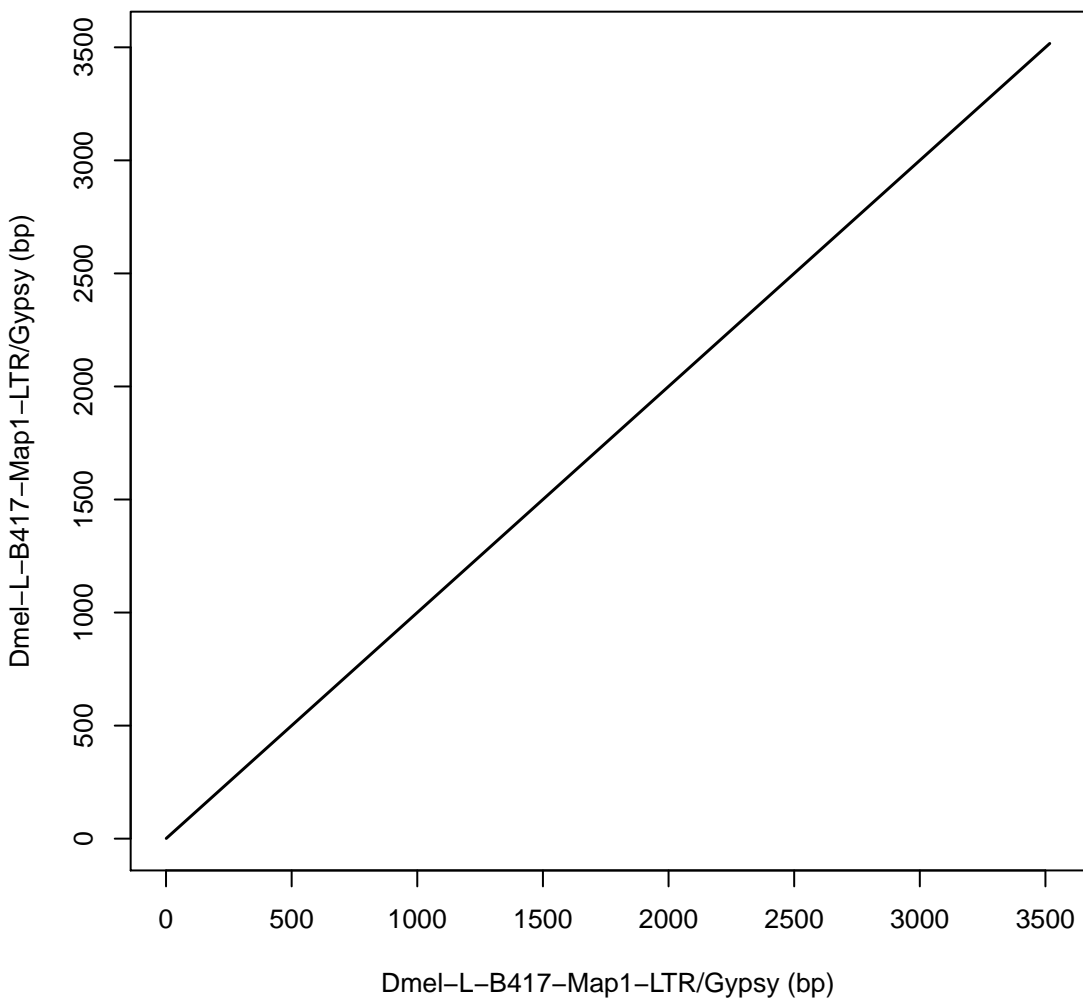
TE: Dmel-L-B417-Map1-LTR/Gypsy
consensus size: 3517bp; fragments: 134; full length: 0 (>=3165.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

