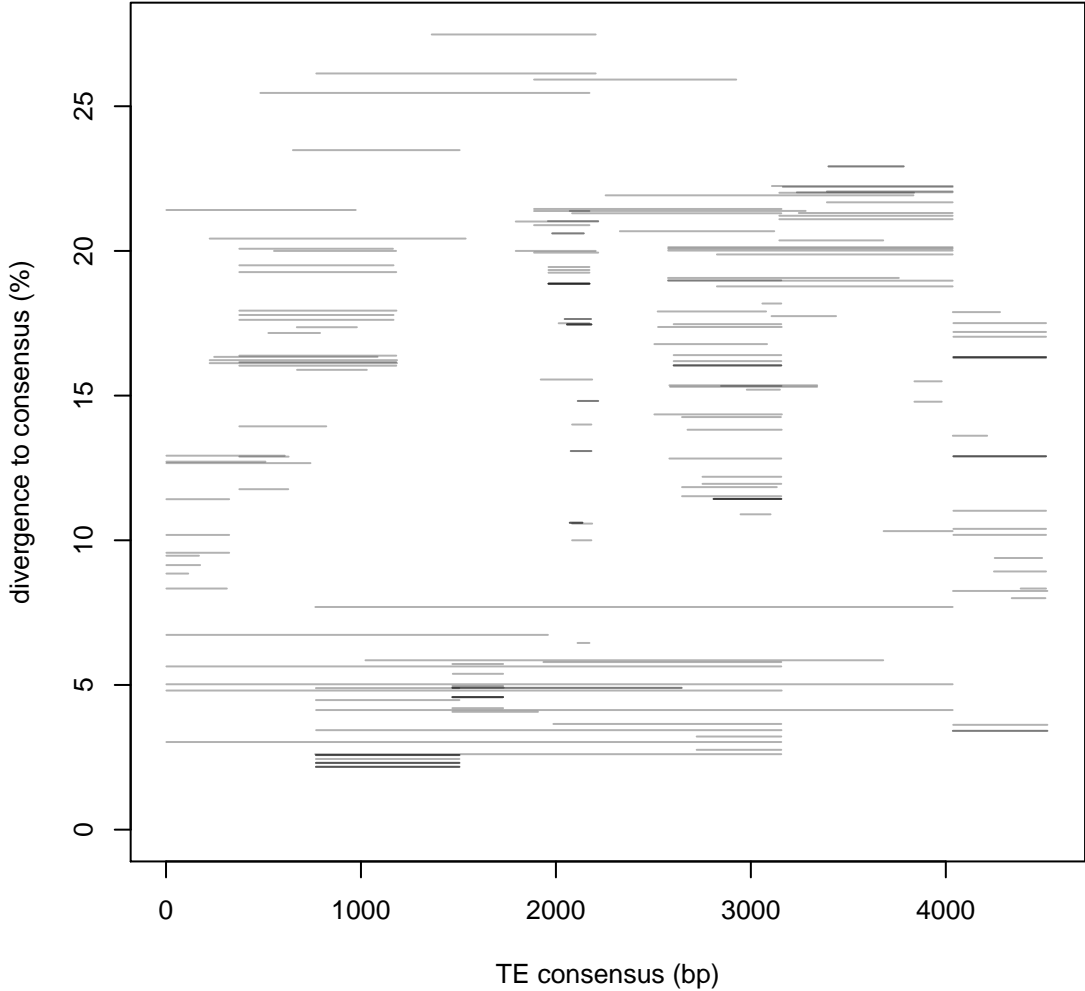
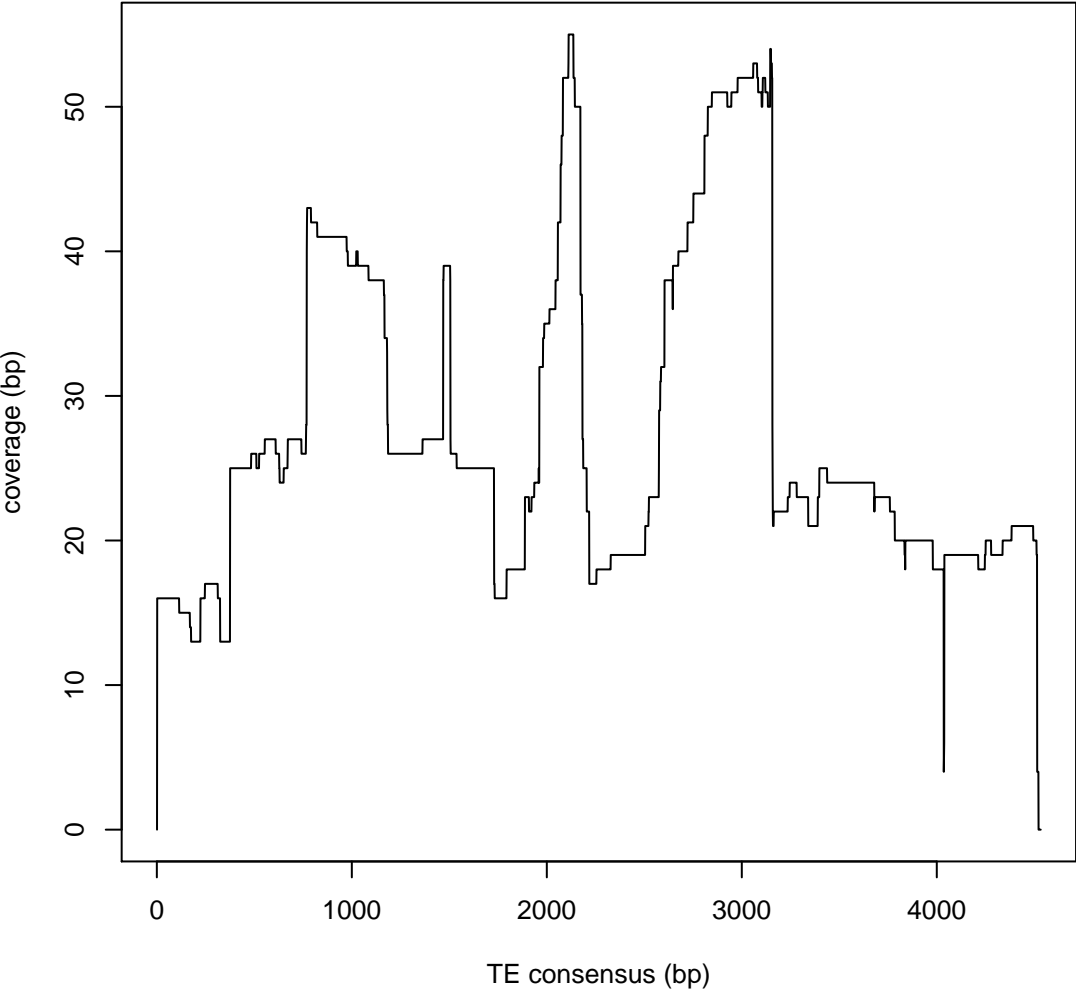


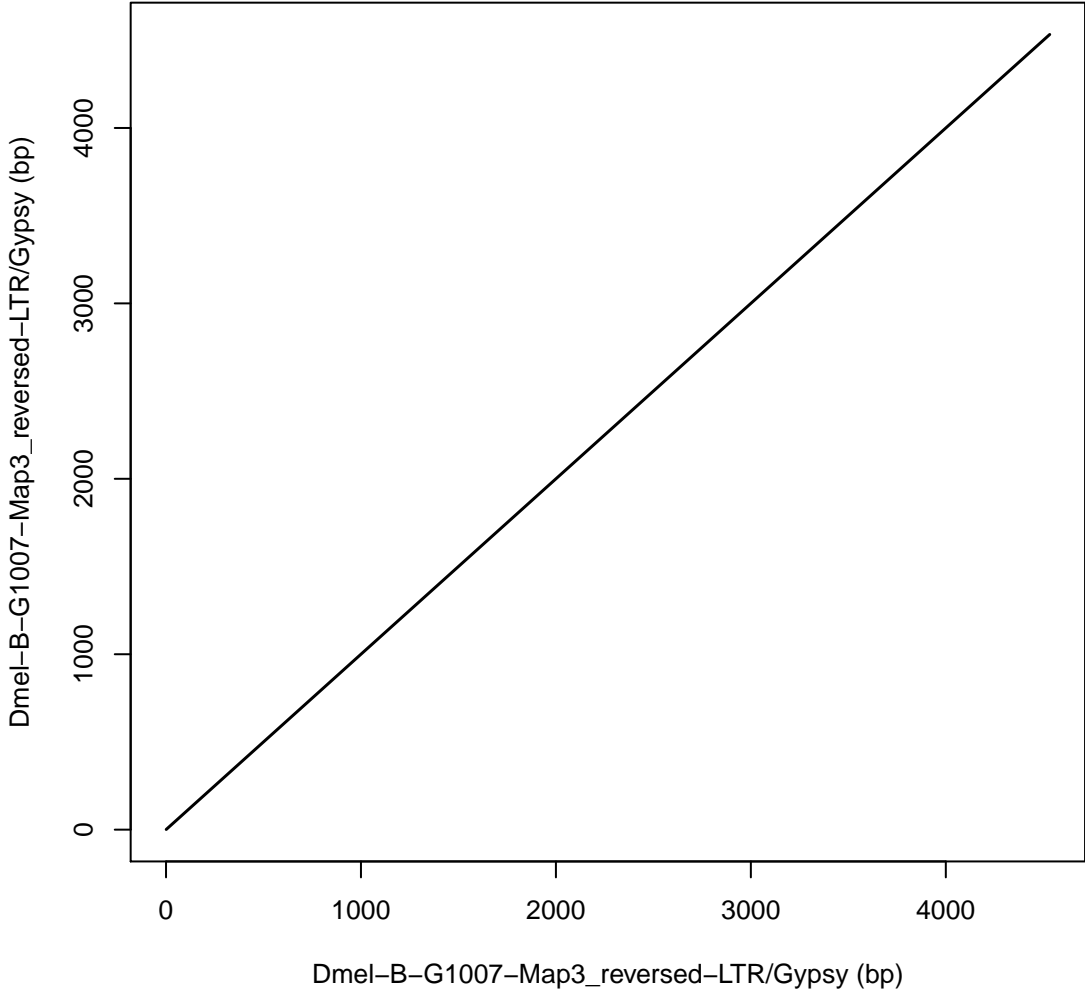
TE: Dmel-B-G1007-Map3_reversed-LTR/Gypsy
consensus size: 4533bp; fragments: 188; full length: 0 (>=4079.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

