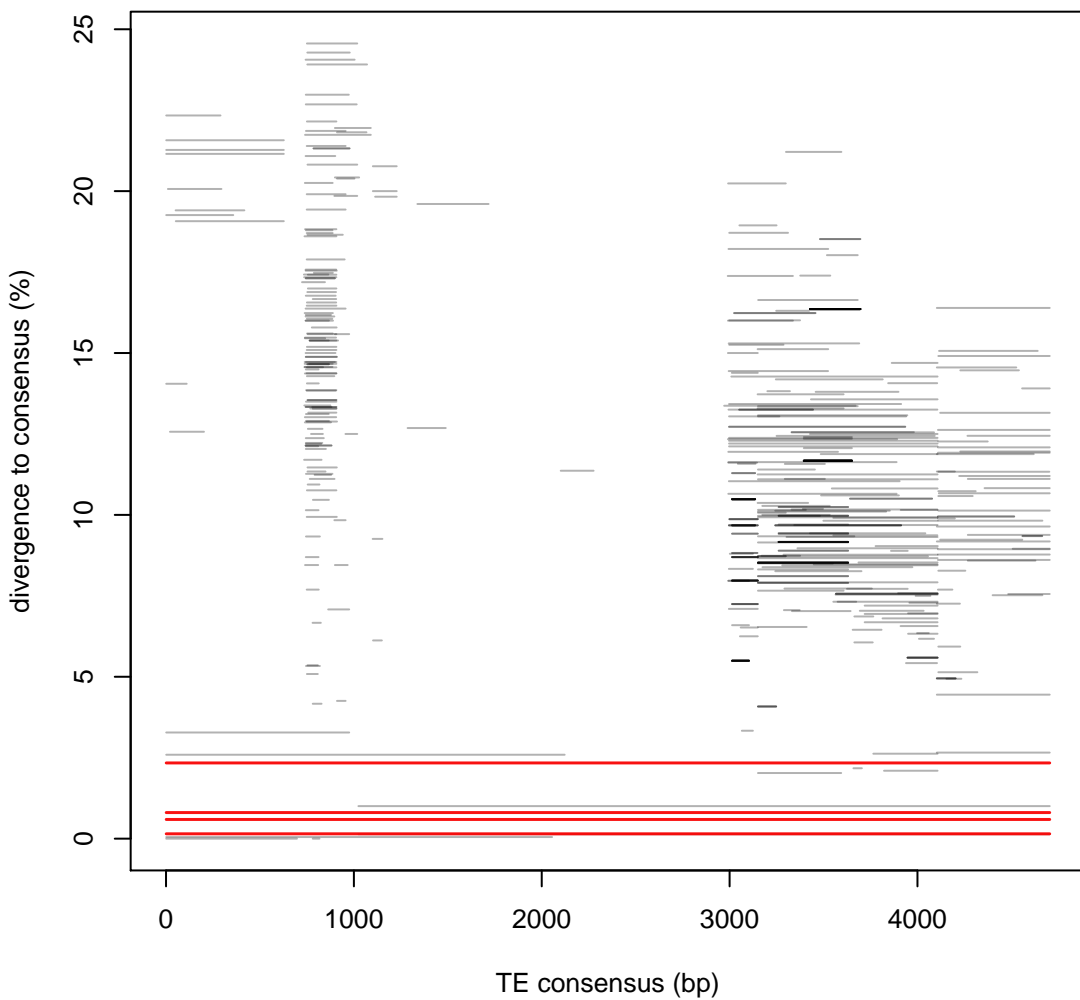
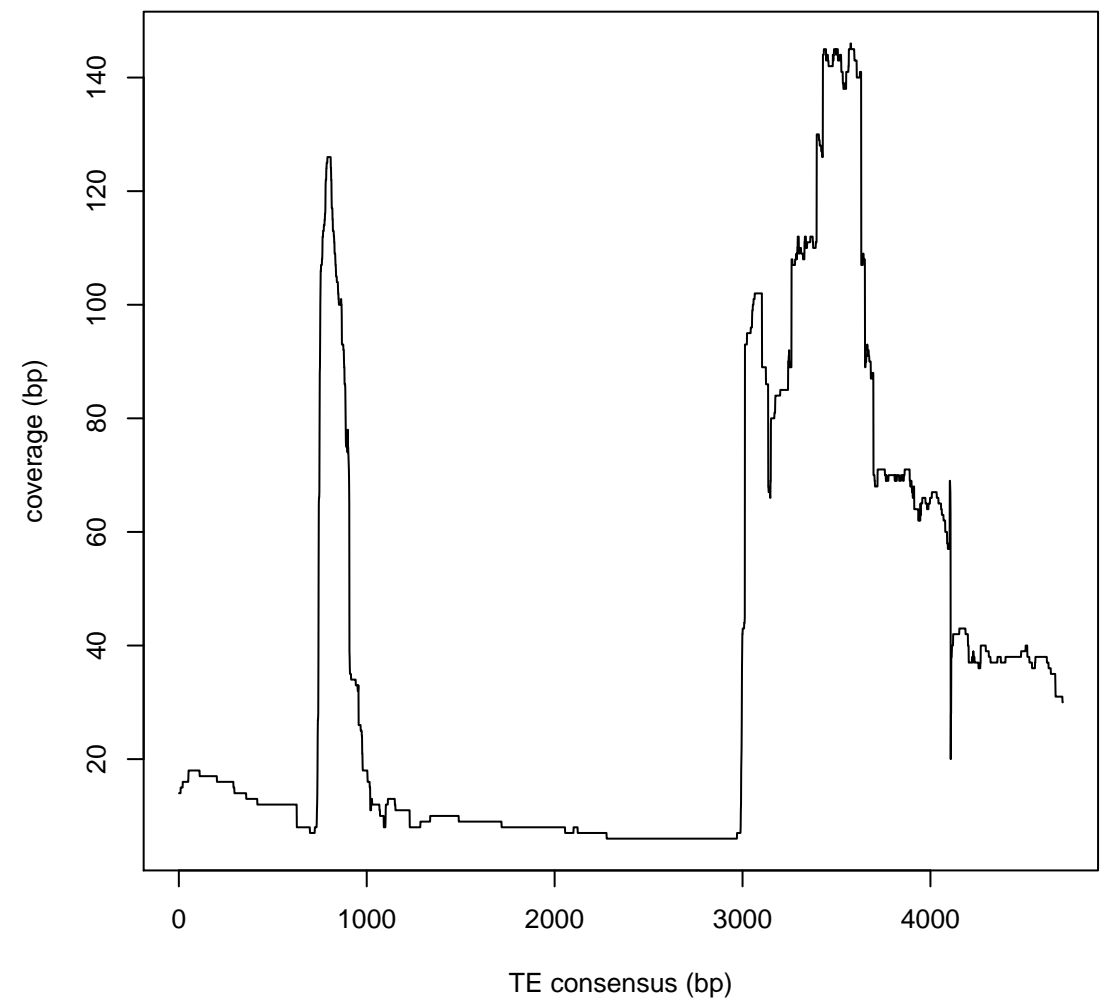


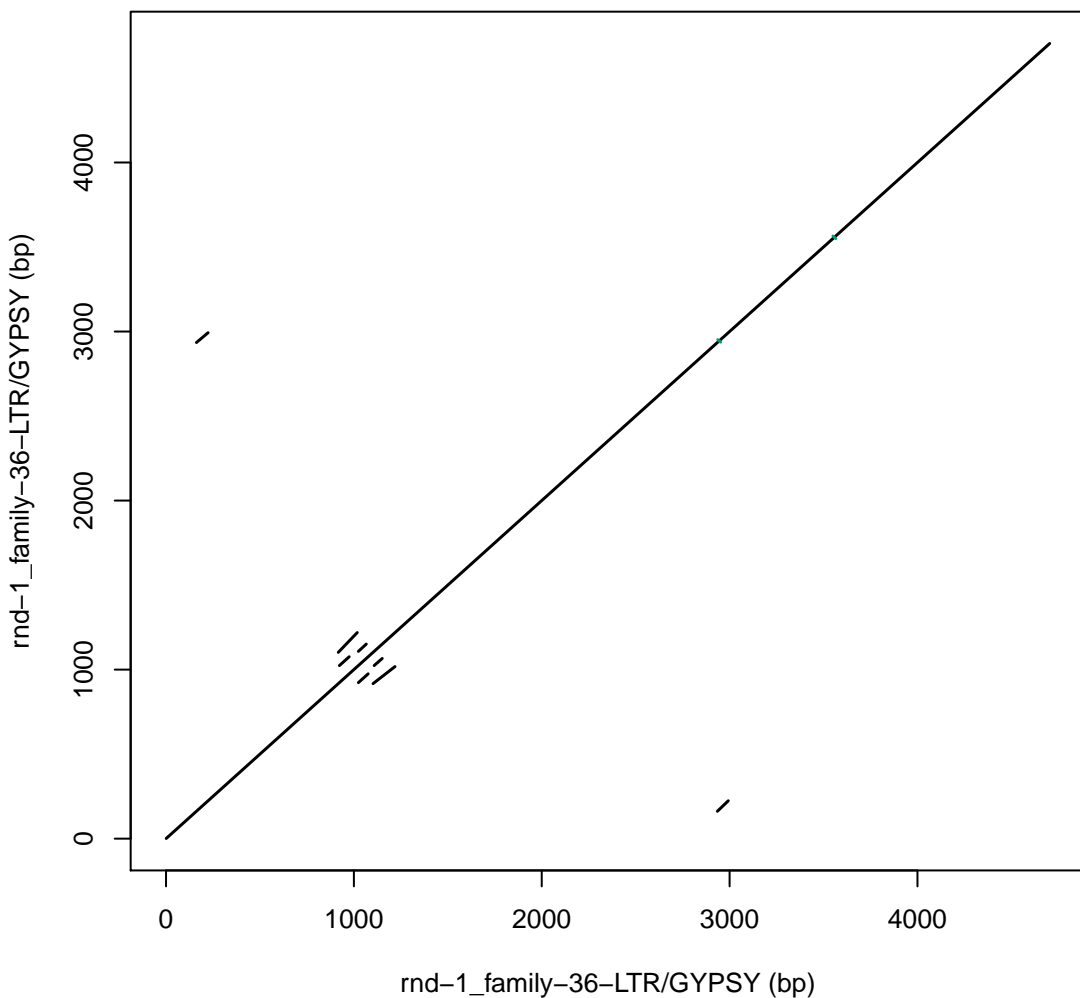
TE: rnd-1_family-36-LTR/GYPsy
consensus size: 4704bp; fragments: 485; full length: 4 (>=4233.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

