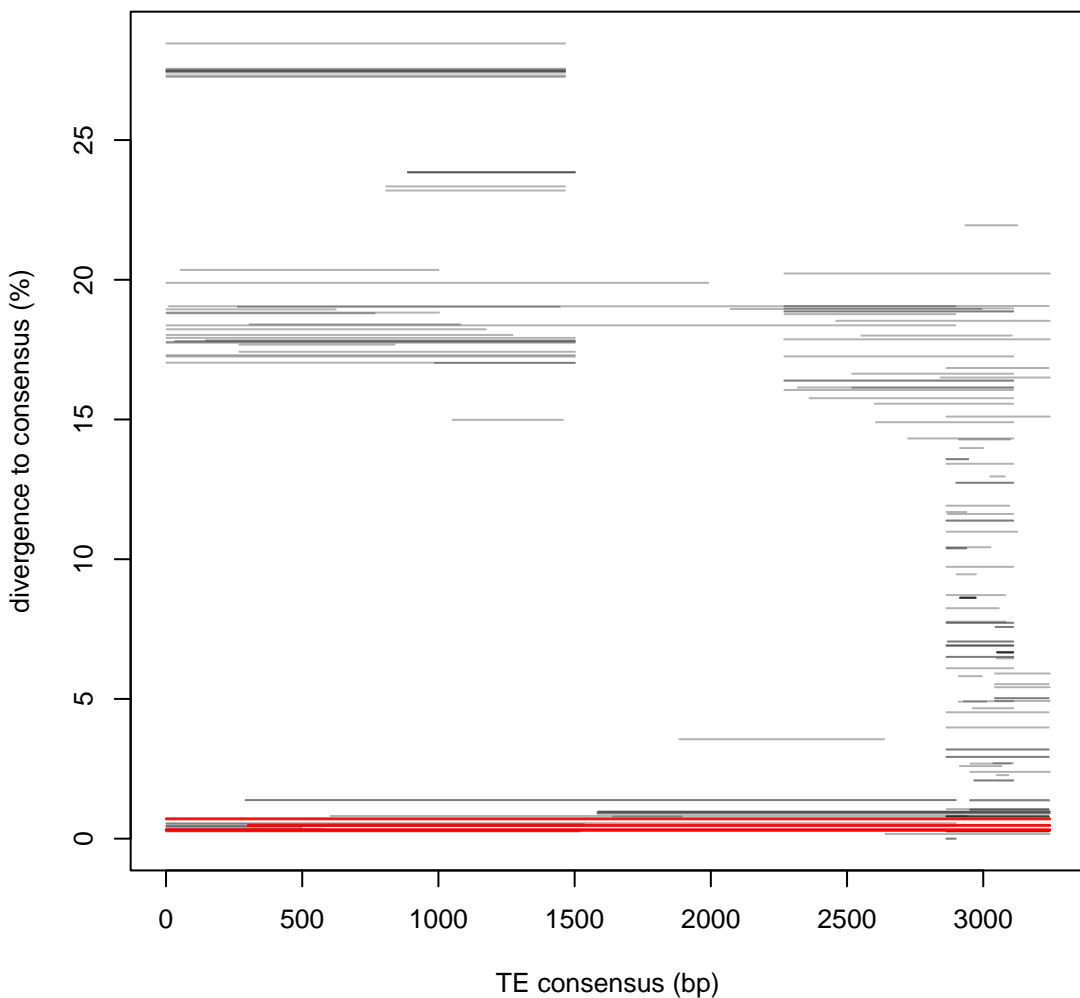
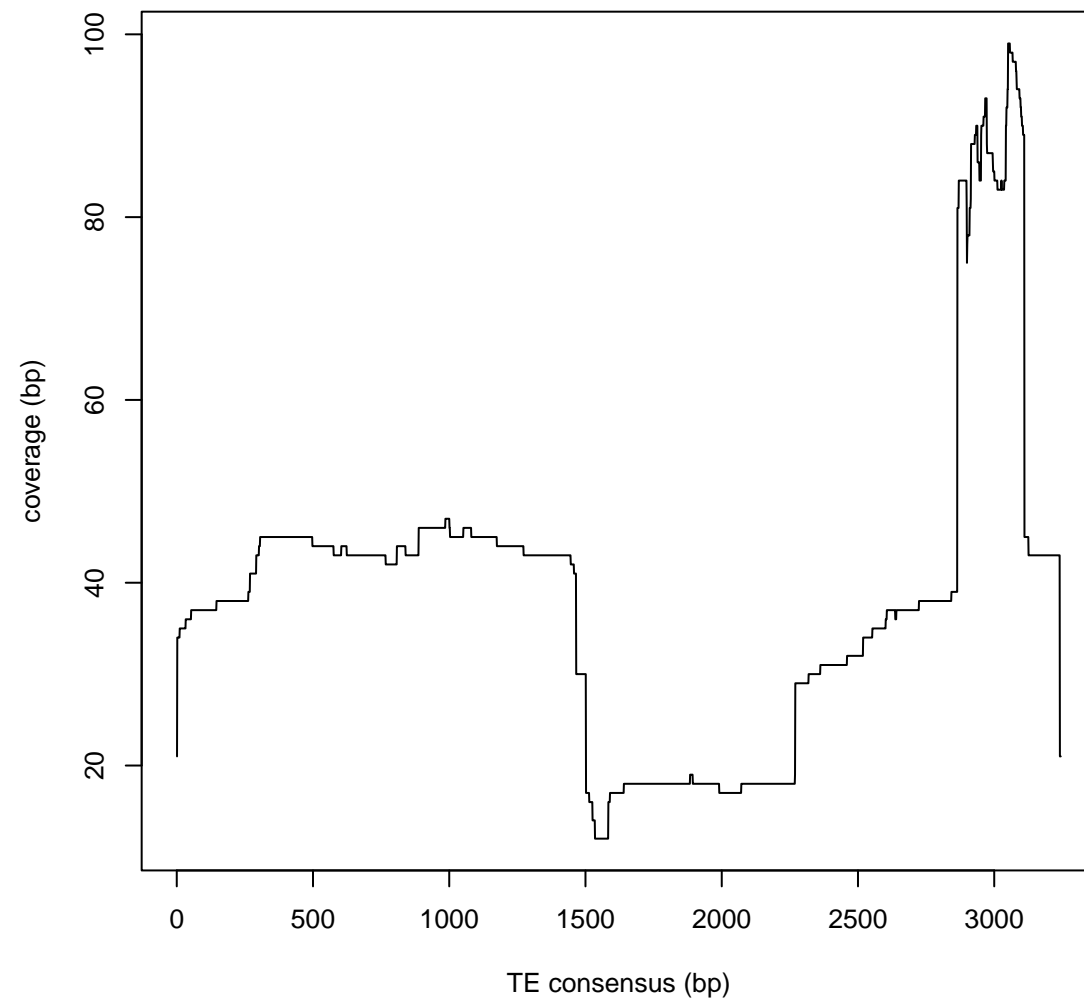


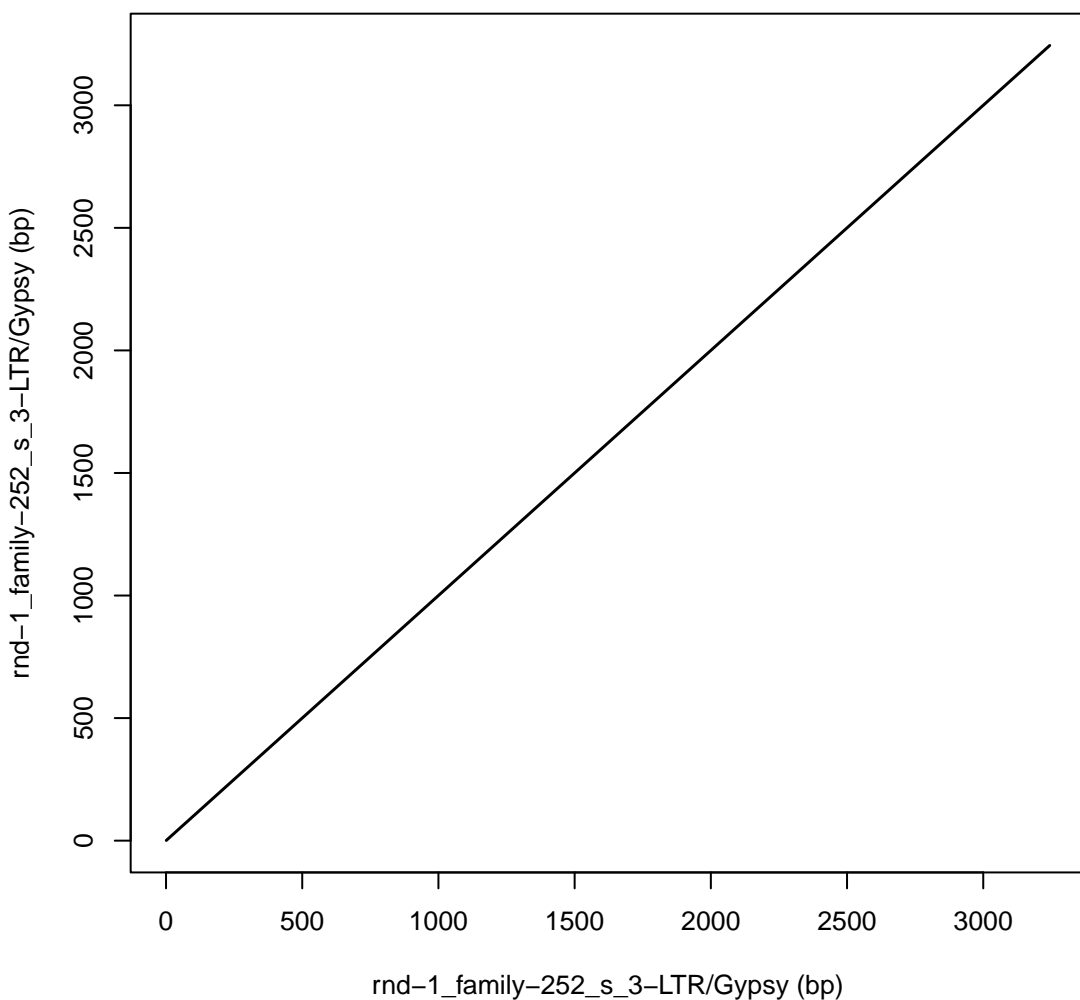
TE: rnd-1_family-252_s_3-LTR/Gypsy
consensus size: 3244bp; fragments: 169; full length: 4 (>=2919.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

